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Maximum Match 100%
Listing first 45 mummaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                        Prad. No. 1s the number of results predicted by chance to have a 
score greater than or equal to the score of the result being printed, 
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                     Score
     A Connected, 101002;

1. / ISINS / Openies / Openies / Openies (Proceed) - MAI, 196 JAN;

1. / ISINS / Openies / Openies / Openies (Proceed) - MAI, 196 JAN;

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(c) 1993 - 2002 Compugen Ltd.
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                               Claim 6; Page 15; 24pp; German
                                                                                                                                                            WPI; 2000-182338/16.
                                                                                                                                                                                                           Maurer K,
                                                                                                                                                                                                                                                                                                                                                              15-JUL-1998;
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## ALIGNMENTS

AAY51581 standard; peptide; 20 AA

Pineapple bromelain protesse peptide fragment from fraction 4 Bromelain, pineapple; protease; blood coagulation; inhibitor; plasmin.

98WO-EP04406

15-JUL-1998; 98WO-EP04406.

(URSA-) URSAPHARM ARZNEIMITTEL CMBH. Grabowska E, Eschmann K;

Composition for inhibiting blood coagulation more efficiently than crude bromelain powder comprises bromelain proteases -

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"fileapple from bronelain processe is used in the treatment or prophylaxis of conditions which are dependent on cyclic nucleocide and other intracellular and/or extracellular purhways. The diseases thou camers, intlamentor, a Abstroceleous and bacterial infections such as choices, diphylaria and pertuasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Porteilabe, pingple, treatment, prevention; prophylaxis, tumour; acid tumour; non-cold tumour; verus induced tumour; non-cold tumour; verus induced tumour; non-cold tumour; language through immunocideficionry virus; maintrition; trauma; burn, buman immunocificionry virus; maintrition; trauma; burn, surgery; generic abnormality; diaberes; mitric oxide; NO; antimicrobial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of purified stem bromelain procease - to treat conditions dependent on cyclic nucleotide pathways e.g. cholera, diphtheria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 108; DB 16; Length 213;
100.0%; Pred, No. 1.8e-10;
tive 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  710 proteinase from pineapple stem.
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                                                                                                                                                                                                                                                                                                                                                                                             Ananas comosus
                                                                                                                                                                                                                                                                                                                                              inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-1994;
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This invention describes a movel use of brossalist processes (I) for imbiliting blood compaintum. If stimulize formation of plasmin, but imbiliting blood compaintum. If stimulize formation of plasmin, but imbilities but formation of fifteen and admission of thromsopytes to montholist, calls. (I) are about 10 times more extraction bromsalist promeins to formation of the most of the formation of the most of the impossible formation and the most operated in the method of the impossible impairs operated.
                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzyme, esp. bromelain, contg. compsn. for control or diarrhoea -
esp. when caused by heat-labile toXIn-producing B. coli, also
inhibite Blud secretion in the gut and promoces absorption
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                                                                                                                                                                                                                          100.0%; Score 108; DB 21; Length 20; 100.0%; Pred. No. 1.1e-11; tive 0; Mismatches 0; Indels 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR47550 standard: protein: 213 AA.
                                                                                                                                                                                                                                                                                                                       1 VPQSIDWRDYGAVTSVKNQN 20
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es 20; Conservative
                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stem bromelain protease.
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                                                                                                                                                                                                                                                      Local Similarity
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Gaps

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Bromelain; protease; proteolytic enzyme; cholera; diphtheria; cancer;
                                                                                                                                                   AAR68974 standard, protein, 213 AA
                                                                                                                                                                                                                                                                                                      Pineapple stem bromelain protease
2 VPQSIDWRDYGAVTSVRQN 21
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Best Local :
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     WUZUUU14253-A1
                                                                                                       oblid tumour; non-solid tumour; virus induced tumour; semunostimulant; immunoseficiency; HTV; selection; malorial malnutrition; trauma; burn; maloria; maloria; malnutrition; trauma; burn; unique; genetic abnormality; diabetes; nitric oxide; NO; antimicrobial.
                                                                                                                                                                                                                                                                                           D385848 proteinase from pineapple stem.
                                                                                                                                                                                                                                                                                                                                                       04-000-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY83107 standard; Protein; 324 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 58-59; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stem bromelain-like proteinases useful for the prevention and treatment of cancer and diseases or conditions which respond to increased nitric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; AAZ93389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CORT-) CORTECS UK LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-MAR-2000
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                                                      Allands company
                                                                                                                                                                                                                                     Proteinase; pinapple; treatment; prevention; prophylaxis; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96 VPQSIDWRDYGAVNEVXOON 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 VPOSIDWRDYGAVISVKNON 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-256998/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       311 AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Crossett B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                       (tirst entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98GB-0019138
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91.7%; Score 99; DB 21
90.0%; Pred. No. 1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 311
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AAY83104
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of camer and diseases or conditions which respond to increased nitrio
oxide production are encoded by nucleic acid sequences isolated from a
pineapple stem.
                                                                                                                                                                                                               immanostimolant; immunodeficiency, HIV;
human immunodeficiency virus, malakria, malnutrition; trauma; burn
surgery, genetic abnormality; diabetes; nitric oxide; NO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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the same activity as a stem bromelain-like proteinase, CCX2.
The proteinases can be used in human or veterinary medicine and
can be used for treating or preventing cancer, e.g. polid tumours
non-polid tumours and virus induced tumours. The proteinases can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 64-66; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AA293391
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                                16-MAR-2000
                                                                                                                                                                                                                                                                                           Proteinase; pinapple; treatment; prevention; prophylaxis; tumour; solid tumour; non-solid tumour; virus induced tumour;
                                                                                                                                                                                                                                                                                                                                                                             674 proteinase from pineapple stem
                                                                                                                                                                                                                                                                                                                                                                                                                                         04-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY83104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY83104 standard; Protein; 351 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         324 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Indels
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The propern designess a frestly of Exemals, the segment of Co. With the Co. With th
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                                                                                                                                                                                                  New isolated component of bromelain - used for treating e.g. touchmanne diseases, transplant rejection, allergic reactions or cancer toxic shock, apoptosis, parasite or pathogen infections or cancer.
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94.7%; Pred. No. 8.8e-10;
tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                    Disclosure, Page 24; 56pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              569 proteinase from pineapple stem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY83098 standard: Protein: 186 AA
                                                                                           Peek K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-GB02784.
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                                                                                     Engwerda C, Mynott TL.
                            (CORT-) CORTECS UK LID.
                                                                                                                                              WPT: 1998-481194/41
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                                                                                                            Stem bromelain-like proteinases useful for the prevention and treatment of carce and diseases or conditions within respond to increased nitric oxide production are encoded by nucleic acid sequences isolated from a
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90.0%; Pred. No. 1.1e-08;
tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 55-57; 72pp; Bnglish,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          123 VPQSIDWRDYGAVNEVKNON 142
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97GB-0003850.
97GB-0004252.
24-AUG-1999; 99WO-GB02784.
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                                                                                                                                                                     Mynott TL, Crossett B;
                                                                                                            (CORT-) CORTECS UK LTD.
                                                                                                                                                                                                                              WPI: 2000-256998/22.
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                                                                                                                                                                                                                                                                                                                                                                                                    pineapple stem -
                                                                                                                                                                                                                                                     N-PSDB; AAZ93388
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25-PEB-1997;
28-PEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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of cancer and diseases or conditions which respond to increased mitric
oxide production are encoded by mucleic acid sequences isolated from a
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02-SEP-1998;
                                                                             24-AUG-1999;
                                                                                                                                                                            16-MAR-2000
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human immunodeficiency virus; malaria; malnurrition; trauma; burn;
surgery, genetic abnormality; diabetes; nitric oxide; NO;
surgery, genetic abnormality; diabetes; nitric oxide; NO;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY83101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY83101 standard; Protein; 186 AA.
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17; Conserv
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98GB-0019138
                                                                                      99WU-GBUZ784
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                                                                                                                                                                                                                                                                                                                                       /note= "Unidentified amino acid"
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Pred. No. 2.7e-08;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
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Best Local Similarity 89.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4, Page 50-51, 72pp; English
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                                                   (CORT-) CORTECS OR LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protestames, pitappie, treatment, prevention; prophylaxis, tumour, solid tumour, one-solid tumour; varus induced tumour; consend timour, varus induced tumour; samunostimulant, immunodeficiency HTV, human immunodeficiency varus, malaria; malnutrition; trauma, burn; surgery, genetic abhormality; diabetes, nitric oxide; NO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         571 proteinase from pineapple stem.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequences encoding proteinases from pineapple stem have the same activity as a stem bromelain-like proteinane, CCX2. The proteinases can be used in human or veterinary medicane and can be used for treating or preventing acancer, e.g. solid tumours more-solid tumours and virus induced tumours. The proteinases can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAZ93385
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Pred. No. 2.7e-08;
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Stem bromelain-like proteinases useful for the prevention and treatment of cancer and diseases or conditions which respond to intro-seased nitrix oxide production are encoded by nuclaic acid sequences isolated from a
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Immunostimulant; immunodeficiency virus; malaria; malnutrition; trauma; burn;
sturgery; genetic absormality; diabetes; nitric oxide; NO;
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89.5%; Pred. No. 2.7e-08;
tive 0; Mismatches 2; Indels
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88 17; Conservative
                                                 Mynott TL, Crossett B;
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                                                                                                                                                     MPI; 2000-256998/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The approximate monthly processing of principle reme hower the approximate of the processing of principle reme hower the processing of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 O, Gabs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Proctatage, plapple, tractment prevention, prophylaxie; tumour; nolid menuoni simon, tractment prevention, prophylaxie; tumour; manuoni simon, ismunoni simon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.0%; Score 95; DB 21; Length 187; 89.5%; Pred, No. 2.7e-08; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Unidentified amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 46-47; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY83100 standard; Protein; 188 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 572 proteinase from pineapple stem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-GB02784.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          POSTDWRDYGAVNEVENDN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17, Conservative
         Crossett B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CORT-) CORTECS UK LID.
                                                                                                   WPI: 2000-256998/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   187 AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference 72
                                                                                                                                                     N-PSDR. AAZ93383
                                                                                                                                                                                                                                                                                                                                                                                                    pineapple stem -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ananas comosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W0200014253-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-SEP-1998;
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Mynott TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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Matches

AAYB3106

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local .
                                                                                                                                                                                                                 25-FEB-1997;
25-FEB-1997;
28-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stem bromelain-like proceinases useful for the prevention and treatment of canner and diseases or conditions which respond to increased nitric oxide production are encoded by nucleic acid sequences isolated from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           graft rejection; antiallergic; antiapoptosis; toxic shock;
antiparasitic; antipathogenic; anticancer; antitumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 61; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pineappie stem -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAZ93390
                                                         WPI; 1998-481194/41
                                                                                                        Engwerda C.
                                                                                                                                                                                                                                                                                                                                                    25-FEB-1998,
                                                                                                                                                                                                                                                                                                                                                                                                        03-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                             MO9838291-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ananas comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ananain; bromelain; CCS; autoimmune; immunosuppressant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ananain N-cerminal sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW80801 standard; peptide; 20 AA
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                                                                                                                                                                                                                                                                                                25-MAR-1997;
                                                                                                                                                             (CORT-) CONTECS UK LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 PQSIDWRDYGAVTSVKNQN 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POSIDWRDYGAVNEVKNON 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                190 AA
                                                                                                        Mynott TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Ilrst entry)
                                                                                                                                                                                                                 97GB-0006119.
97GB-0003827.
97GB-0003850.
97GB-0004252.
                                                                                                                                                                                                                                                                                                                                                    ARMO-CROOPAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89.54;
                                                                                                        Peex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No.
                                                                                                        Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.8e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 190;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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ID AAYS
XX
AC AAYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The parent disciouse a fraction of bromabilin, designated CGs, which collisions between the second collisions and the collisions of 27.8 D) as described by 200 Polisions. (i) and described political of 27.8 D) as described by 200 Political collisions are collisions as a collision of 27.8 D) as described by 200 Political collisions. (ii) an apparent political collisions are collisions and collisions are collisions and collisions are collisions. (iii) aspecting the components by fast the collisions are collisions and collisions are collisions. (iii) aspecting the collisions are collisions and collisions are collisions. (iii) aspecting the collisions are collisions and collisions are collisions. (iii) aspecting the production collisions are collisions are collisions. (iii) aspecting the production collisions are collisions. (iii) aspecting the production collisions are collisions. (iii) aspecting the production of growth factors and collisions are collisions. (iii) the collisions are collisions. (i
     RESULT 14
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Best Local Similarity
This invention describes a novel use of bromelain proteases (I) for inhibiting blood coagulation. (I) stimulate formation of plasmin, but
                                                                              Claim 8; Page 16; 24pp; German
                                                                                                                                  Composition for inhibiting blood coagulation more efficiently than crude bromelain powder comprises bromelain proteases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pineapple bromelain protease peptide fragment from fraction 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY51582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY51582 standard; peptide; 20 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   toxic shock, apoptosis, parasite or pathogen infections or cancer
                                                                                                                                                                                                                 WPI; 2000-182338/16
                                                                                                                                                                                                                                                                     Maurer K,
                                                                                                                                                                                                                                                                                                                    (URSA-) URSAPHARM ARZNEIMITTEL GMBH.
                                                                                                                                                                                                                                                                                                                                                                              15-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ananas comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bromelain; pineapple; protease; blood coagulation; inhibitor; plasmin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       000Z-XW-RT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune diseases, transplant rejection, allergic reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 VPQSIDWRDYGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VPQSIDWRDSGAVTSVKNQ 19
                                                                                                                                                                                                                                                                     Eckert K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                              98WO-EP04406
                                                                                                                                                                                                                                                                                                                                                                                                                                  98WC-EPU4406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94.7%;
                                                                                                                                                                                                                                                                     Grabowska E, Eschmann K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 93; DB 19;
Pred. No. 4.4e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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Length 20 Indels 0 Gaps

New isolated component of bromelain - used for treating e.g.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bromelain, CCS(P2, BRK-2 phosphorylation; interleukin-2; offer cell proliferation; interpretent profits at the prostine; onner; parasite, pathagan inferction; oproxine; account parasite, pathagan inferction; interpretent, accommand takeses; agar for lefection; transplant rejection; allergia resection; taxon account or processing the properties.
inhibit both formation of fibrin and adhesion of thrombocytes to controlled and the state of the condition of the condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Browelain fractions useful in the prevention and treatment of cancer, pressives and pathogen infections and autoimmune diseases comprise ppressive such biological activity.
                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                        86.1%; Score 93; DB 21; Length 20; 94.7%; Pred. No. 4.4e-09; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-terminal peptide from a 26 kDa protein of bromelain.
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Matches 18; Conserv
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or transplant rejection by a host, allergic reactions, toxic shock or apoptosis. The fractions and proceins may also be used to prepare agents for the treatment of the above diseases and conditions.
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                                                                                                86.1%; Score 93; DB 21; Length 21; 90.0%; Pred. No. 4.6e-09;
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## ALIGNMENTS

standard; peptide; 20 AA.

antiparasitic; antipathogenic; anticancer; antitumour.	antiparasitic;
graft rejection; antiallergic; antiapoptosis; toxic shock;	graft rejection
Ananain; bromelain; CCS; autoimmune; immunosuppressant;	Ananain; brome

WPI; 1998-481194/41.	Engwerda C, M	(CORT-) CORTECS UK LTD.	03-SEP-1998; 25-FEB-1998; 25-FEB-1997; 25-FEB-1997; 28-FEB-1997; 28-FEB-1997;	
94/41	Mynott TL,	SUK	98WO 97G8 97G8 97G8 97G8	
	Ħ	LTD.	98WO-GB00590. 97G8-0006119. 97GB-0003827. 97GB-0003850. 97GB-0004252	
	Peek K,		190	
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Page 2

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Gaps

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The present expense represents a Vicential sequence from a fraction of broadcain, designated CGS/P2, comprising proteins of molecular weights for which are sequenced to the properties of molecular weights for when measured by shame sequenced. The pecklose block bits of the protein of the pr
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CO++ T cell proliferation; mitogan activace procent kinase; MAP kinase;

growth factor; cytokine; concer, parasite, pathogen infection;

faminopupressant; actionment effects; cytokine; Capiction;

Cramaplant rejection; allegist reaction; toxic shock; apoptosis.
infinititing ploto compainted. (I) stillable formation of plassin, but
infinite both formation of fibrin and adhesion of thrombopyses to
endothalia (2018. II) rest about 10 tiens more active then bromedain
crude powder, 1.e. comparable with grappockinses. AMYSISSI and AMYSISS
represent Powder, and the present of the properties of the properties of the properties of the properties of the present indicate of the present of t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A-terminal peptide from a fraction of bromelain designated CCS/P2.
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100.0%; Pred. No. 1.6e-10;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                  Sest Local Similarity
                                                                                                                                                                                                                                                        20 AA;
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                                                                                                                                                                                                                                                               Seguence
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                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition for inhibiting blood coagulation more efficiently than crude bromelain powder comprises bromelain protesses
       New isolated component of bromelain - used for treating e.g. 
uncommune diseases, transplant refection, altergic reactions or toxic shock, apoptosis, parasite or pathogen infections or cancer
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100.0%; Pred. No. 1.6e-10;
ive 0; Mismatches 0;
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                                                                                                                                                    Disclosure, Page 24; 56pp; English
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RESULT 4
AAY84391
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The present exquence trapresents a N-terminal sequence trom a traction of bromelain, designated CCP/27, comprising processing to seasonable way 2, 24 ya Nb amanya 6 km 2, 24 
                                                                                                                                                                      Bromelain fractions useful in the prevention and treatment of cancer 
parasites and pathogen infections and autoimmune diseases comprise 
proteins with biological activity -
                                                                                                                                                                                                                                                                                                                                                                                                                                              16-MAR-2000
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CO4+T cell proliferation; mitogen activated protein kinase; MAP kinase;
growth factor; cytokine; cancer; parasite; pathogen infection;
immunosuproseant; autoimune, disases; graft rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-terminal peptide from a fraction of bromelain designated CCS/P2
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                                                                                                                                   Claim I; Page 51; 89pp; English.
                                                                                                                                                                                                                                                        WPI; 2000-256999/22
                                                                                                                                                                                                                                                                                           Mynott TL, Peek K,
                                                                                                                                                                                                                                                                                                                             (CORT-) CORTECS UK LTD.
                                                                                                                                                                                                                                                                                                                                                                                                           24-AUG-1999;
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Pred. No. 2.3e-10;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                        Mynott TL,
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The broad-inf fractions and proceins are used in sedicine as they inhibit the misons set treed process. They have been set to set the control of the misons set treed process. They have been set to set the control of the set of particles of particles in factorism. They can be used as an amount of the control of the set of particles of particl

100.0%; Score 105; DB 2 Pred. No. 3e-10; DB 21; Length

0

Indels 0 Gaps 0,

VPQSIDWRDSGAVTSVKNQG 20 0 Mismacches

AAY84394 standard; peptide; 21 Š

(first entry)

peptide from a 24 kDa protein of bromelain.

Bromelain; CCS/92; SRK-2 phosphorylation; interleukin-2; CCG+ To call proliferation; mitogen activated protein kinase; MAP kinase; gowth factor; cyrokina; cancer; parasite; pathogen infection; immunospessant; autolimunos disease; graft re-phogen infection; transplant rejection; allegge reaction; coak shock; apoptosis.

/noces Location/Qualifiers "not specified"

98GB-0019137 99WO-GB02792.

(CORT-) CORTECS OK LID

Peek K, Tyreman DR;

Bromelain fractions useful in the prevention and treatment of cancer, parasites and pathogen infections and autoimmune diseases comprise proteins with biological activity.

Claim 3; Page 52; 69pp; English

The present sequence represents a N-cemical sequence from a 24-bb process in a measured by SER-PGI. The process is desired from a fraction process in a measured by SER-PGI. The process is desired from a fraction 31-3-bbs and 36-6 bbs when measured by SER-PGIC or 21-4-59 bbs and 21-3-59 parasites or pathogen infections. They can be used as an

proliferation

8×99999

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infections, or cancer. The CCS fraction is postulated to consist of the yequeine protesses strains and compania, together with various other components. The present sequence represents the N-terminal of compasin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bromelain, pineapple, protease; blood coagulation, inhibitor, plasmin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pineapple bromelain procease peptide fragment from fraction 4.
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                                                                                                                                                                                                    89.5%; Score 94; DB 19; Length 20; 90.0%; Pred. No. 1.1e-08; ive 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                 1 VPQSIDWRDSGAVTSVKNQG 20
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es 18; Conservative
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Matches 18, Conserv
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AC AAR4
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immunosupressant for treating or preventing subclommune diseases, graft or treating the rejection by a host, allergic reactions, toxic shock or soppores; The frottions and proteins may also the used to prepare agents for the above diseases and conditions.
                                                                                                                                                                                                                                                                                                                     Gabs
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                                                                                                                                                                                                                                               95.2%; Score 100; DB 21; Length 21; 95.0%; Pred. No. 1.1e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Comosain, bromelain, CCS, autoimmune; immunosuppressant;
graft rejection, antiallargic, antiappicells, toxic shock,
antipassatzic, antipathogenic; anticancer; antitumour.
                                                                                                                                                                                                                                                                                                              1: Indels
                                                                                                                                                                                                                                                                                                                     0: Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                    1 VPOSIDWRDSGAVTSVKNOG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VPOSTDWRPSGAVTSXKNOG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Comosain N-terminal sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-GB00590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97GB-0006119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97GB-0003850.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CORT-) CORTECS UK LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-481194/41,
                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ananas comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W09838291-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Engwerda C,
                                                                                                                                                                                                                                                                                                                     .61
                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW80802;
                                                                                                                                                                                                                                                      Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 6
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                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence is that of seem bromelain processe, a mixture of various modelies derived from the sees of the phrappie. The process is a cyterious processe used to treat infectious starrhoes processe in a cyterious continuous conditions of the processe which is the processe that the interface of the totals and prevents attachment of bacteria to the interestical lining. It also minities serverious couled by secretagopues, e.g. PGES, thoughlylline, etc. and has no adverse street municipal minities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 11-13, 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzyme, esp. bromelain, contg. compsn. for control or diarrhoea esp. when caused by heat-labile toxin-producing E. coli, also inhibits fluid secretion in the gut and promotes absorption
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUN-1992;
20-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Annus comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gastro-intestinal tract; GI tract; absorption.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Proteolytic enzyme; cysteine protease; diarrhoea; fluid secretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stem bromelain protease
             24-JUN-1994;
                                                                                                                                                          rineapple stem prometain procease.
                                                                                                                                                                                    09-AUG-1995
                                                                                                                                                                                                                                    AAR68974 standard; protein; 213 AA
                                                                                                                                                                                                                                                                                                                                                                                                              sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1994-025889/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mynott TL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CORT-) CORTECS LTD
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                                                                  MC6200168-W
                                                                                          Ananas comosus
                                                                                                                    inflammation.
                                                                                                                               brometain; procease; proceotytic enzyme; chotera; diphtheria; cancer;
                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                            1 VPQSIDWRDSGAVTSVKWQ 19
                                                                                                                                                                                                                                                                                                    VPQSIDWRDYGAVTSVKNQ 20
                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                              213 AA;
                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                    (Ilrec entry)
             94WO-GB01368
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93GB-0008164
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                                                                                                                                                                                                                                                                                                                                                                      94.74
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                                                                                                                                                                                                                                                                                                                                                                      Score 93;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                         2.4e-07;
ches 1;
                                                                                                                                                                                                                                                                                                                                                                                   DB 15;
                                                                                                                                                                                                                                                                                                                                                                                 Length 213;
                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 10
AAY84395
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Macches
                                                                                                                                                                                                                                                                                                                                                         Bromelain; CGS/P2; BRK-2 phosphorylation; interleukin-2; Bromelain; CGS/P2; BRK-2 phosphorylation; interleukin-2; MRK-2 phosphorylation; interleukin-2 account parasite; pathogen infection; immunosphessant; autoimmuno diesese; graft rejection; transplant rejection; allergic reaction; coate Shock; apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prophylaxis of conditions which are dependent on cyclic nucleotide and other intracellular and/or extracellular pathways. The disease include cancer, inflammation, atherosci-cirosis and baccerial infections such as cholera, diphtheria and pertussis.
             parasites and pathogen infections and autoimmune diseases comprise proteins with biological activity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pineapple stem bromelain protease is used in the treatment or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 35; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of purified stem bromelain protesse - to treat conditions dependent on cyclic nucleotide pathways e.g. choiers, diphtheria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mynott TL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUN-1993;
                                       Bromelain fractions useful in the prevention and treatment of cancer
                                                                  WPI; 2000-256999/22
                                                                                          Mynott TL, Peex K,
                                                                                                                                               1866T-ASS-ZO
                                                                                                                                                                       24-AUG-1999;
                                                                                                                                                                                                                                                              Misc-difference 10
                                                                                                                                                                                                                                                                                                                                                                                                                                        N-terminal peptide from a 26 kDa protein of bromelain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY84395;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY84395 standard; peptide; 21 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-051751/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CORT-) CORTECS LTD
                                                                                                                    (CORT-) CORTECS UK LTD
                                                                                                                                                                                                 16-MAR-2000
                                                                                                                                                                                                                            WO200014254-A2
                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                 Windlig Comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             213 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                               98GB-0019137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93GB-0013188
                                                                                                                                                                         99WO-GB02792
                                                                                                                                                                                                                                                    /note= "not specified
                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                              - a 2 out /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88.6%;
94.7%;
                                                                                          Tyreman DK.
                                                                                                                                                                                                                                                                              "not specified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 93;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.46-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 16; Length 213
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The proteint (septions represents Neterachia) and sequence from a fraction protein (as measured by 502-A05). The protein laid derived from a fraction to protein it as measured by 502-A05). The protein laid derived from a fraction of formal laid derived from a fraction of the formal laid derived from a fraction of the formal laid of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stem bromelain-like proteinages useful for the prevention and treatment of canners and diseases or conditions which respond to intreaged mitric oxide production are encoded by nucleic acid sequences isolated from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Proceinage plaapple, treatment; prevention; prophylaxis; tumour; solid tumour; non-solid tumour; non-solid tumour; non-solid tumour; immunostimalan; immunosticateny HIV, immunosticateny HIV, superiorismi manutecite; temperiorismi purni, surgery; genetic abnormality, diabetes; nitrio oxide; NO; surgery; genetic abnormality, diabetes; nitrio oxide; NO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequence ancoding proceinage from pineapple stem have 
the proceinage and proceinage from pineapple and 
The proceinages can be used in haman or weretrancy meditains and 
can be used for restain or preventing cancer, so, solid unmorra, 
non-solid unmours and virus induced temours. The proceinages can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81.0%; Score 85; DB 21; Length 21;
85.0%; Pred. No. 3.5e-07;
cive 0; Mismatches 3; Indelu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAYB3105 standard, Protein, 311 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     710 proteinase from pineapple stem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 58-59; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPQSIDWRDSGAVTSVKNQG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPOSIDWRDSGAVTSVXPXG 20
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es 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CORT-) CORTECS UK LID.
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N-PSDB; AAZ93389.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MO200014253-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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              88888888888888888
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                                                               The processing sequence represents a herarizable appearent from a fraction processing sequence represents a herarizable appearent for the processing of measured by SBS-PAGE1. The protein is derived from a fraction of processing of measured by SBS-PAGE1. The proteins is derived from a fraction for the processing of measured for the processing of measured for the processing of measured for measured for measured for measured for the production and CRA-Treat proplets from the production and CRA-Treat production for the production and CRA-Treat production for the production and CRA-Treat Treat for the form of the form of production and craft for the first form of the form of pathogen from fractions and production and craft for the first form of the first form in the form of the first form in the first form i
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COM-T cell proliferation; mitogen activated procent kinase, MAP kinase;
growth factor; Orokoline; cancer; parasite, pathogen infection;
immiropurpessant; autofiname disease; graft or Pojetion;
Crassplant rejection; allergic reaction; toxic shock; apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bromelain fractions useful in the prevention and treatment of cancer, paparasites and pathogen infections and autodemune diseases comprise proteins with biological activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0, Gaps
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89.5%; Pred. No. 1.1e-07;
Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-terminal peptide from a 24 kDa protein of bromelain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note= "not specified"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY84396 standard, peptide, 21 AA.
              Claim 5: Page 52: 69np: English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPOSTDWRDSGAVTSVKNO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-GB02792.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUL-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI: 2000-256999/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ananas comosus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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RESULT 13
AAY83107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       also be used as an immunostimilant or in the preparation of an immunostimilant for treat immunostimilant in training from mainuration, infection (e.g. HIV and salaria), tumours (e.g. hyphotic, systems), transmi (e.g. burne, wounds, surgery) medical preference (e.g. with rough send (e.g. strong disportable). The production of the component immunostimilant in the component immunostimilant in the component immunostimilant in the component immunostimilant in the component of diseases or callia), daletes and old age. The processme can also be used in the treatment or prevention of diseases or conditions which respond to increased nitric oxide (ND) production conditions which respond to increased nitric oxide (ND) productions of the component of an antial croad-all agent or the component of an antia
The acquartee encoding procedures from primagels are have 
from procedures on the used in homeon or vactionary saddiction and 
can be used for creating or preventing cancer, e.g. solid timester 
and solid timester and virtue influence timester. The procedures can 
temporary temporary to the procedures and the procedures of 
immunostruction in facetion (e.g. att your and making), timester e.g. 
whiterittion, infection (e.g. att your and making).
                                                                                                                                                                                                                                                                                                                                                 Stem bromelain-like proteinases useful for the prevention and treatment of cancer and diseases or conditions which respond to increased nitric exide production are encoded by nucleic acid sequences isolated from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunostimilant; immunodeficienty, HIV;
human immunodeficiency virus, malaria; malnutrition; trauma; burn
surgery; genetic abnormality; diabetes; nitric oxide; NO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY83107 standard; Protein; 324 AA
                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                    pineapple stem -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-RADR:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MYNOTE TL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CORT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antimicrobial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Proteinase; pinapple; treatment; prevention; prophylaxis; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D385848 proteinase from pineapple stem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPOSIDWRDSGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-256998/22
                                                                                                                                                                                                                                                                 4; Page 64-66; 72pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ93391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Crossett B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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AAY83104
ID AAY83
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inymphoid, mystems), traums (og.) burns, wounds, surgery) medical treatment (og.) with thougs much as mercuis, cyclospostan and cyclospospanido), procesh loss (og.) and shoutering and burns and burns of the cyclospostan and c
                                                                                                                                                                                                                                                                                                                                                                                                                                 Stem bromelain-like proteinages useful for the prevention and itsetment
of caneer and diseases or conditions which respond to increased intentent
oxide production are encoded by nucleic acid sequences isolated from a
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solid tumpour; non-moild tumpour; virus induced tumpour;
immunostimulant; immunodeficions; HIV;
human immunodeficionsy virus; malaria; malnutrition;
surgery; genetic abnormality; diabetes; nitric oxide;
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     also be used as an immunostimulant or in the preparation of an immunostimulant to rin the preparation of an immunostimulant to test immunostimulant are arising from animulantition, infection (e.g. HIV and malaria), tumours (e.g. malaria), fumbours (e.g. Hivania), publical, surgery) medical, lymphoid, myathon y tramma (e.g. burns younds, surgery) medical, treatment (e.g. with drugs much as steroids, cyclosporin and cyclophosphanids), potenth loss (e.g. in dishortnes and burns),
                                                                                                                                                                           The sequences encoding proceinases from pineapple stem have the same activity as a stem bromelain-like proteinase, CCX2. The proceinases can be used in human or veterinary medictine and can be used for treating or proventing canner, e.g. solid tumours mon-solid tumours and vitue, induced tumours. The proteinases can
                                                                                                                                                                                                                                                                                                                                                                                                          pineapple stem -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAZ93388
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                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 55-57; 72pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000-256998/22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prophylaxis; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 324
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trauma; burn, NO;

0 Gaps

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0; Gabs
equencie, Admonstalities (e.g., in combinand immunosidicinency partients lacking 6 or 7 cells), diabetes and old age. The proteinses can also be used all the treatment or prevention of diseases or conditions which respond to increased mixtic oxide (MOD) production, or as an antimicrobial agent or the component of an antimicrobial agent or the component of an antimicrobial agent.
                                                                                                                      Query March

80.04; Score 84; DB 21; Length 351;
Best Local Statlarity 84.28; Pred. No. 1.3e-02;
Matches 16; Conservative 0; Mismatches 3; Indels
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                        friticum, wheat seed; cysteine proteinase; gluten; baking.
                                                                                                                                                                                                                                                                                                                                                            friticum sp. cysteine proteinase #1.
                                                                                                                                                                                                                                                                             AAW89556 standard; Protein; 377 AA.
                                                                                                                                                                                                     123 VPOSIDWRDYGAVNEVKNO 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98JP-0098140.
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                                                                                                                                                                             1 VPQSIDWRDSGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                 16-MAR-1999 (first entry)
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                                                                                                351 AA:
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Matches
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Gaps

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Query Match 79.04; Score 83; DB 20; Length 377; Deet Local Similarity 70.04; Pred. No. 2.1e-05. Metches 14; Conservative 3; Mismatches 3; Indels

377 AA;

Sequence

Search completed: November 21, 2002, 16:08:39 Job time : 31 secs

138 LPRSVDWRQKGAVTGVKNQG 157

1 VPQSIDWRDSGAVTSVKNOG 20

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The present sequence represents a cysteine proteinase isolated from whether seed (Triftium sp.). The cysteine profelase is useful for improving gluten for use in the bakery process.

New DNA coding cysteine proteinase originating from wheat seed - useful for improving gluten for use in bakery process

Claim 1; Page 15-16; 29pp; Japanese

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200 200 200 200 200 200 200 200 200 200		Pred. No score gr and is d and is d	Post-processing: Database :	Total number of Minimum DB seq Maximum DB seq	Searched:	Sequence: Scoring table:	Title: Perfect score:	protein -	
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Spermatophyta, Magnollophyta, eudicotyledoma, core eudicote; Rosidae,
euroside II; Erassicales, Caricaceae, Carica:
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01-0AN-1997 (Rel. 13, Last emperence update)
18-UWN-2007 (Rel. 14, Last annotation update)
Chymopapain precursor (RC 3.4.22.6) (Rapaya proteinase II) (PPII).
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Matson D.C., Yadrobih M., Dynn K.R.;
"The amino acid sequence of chymopapain from Carica papaya.";
Bjochem. J. 266:75-81(1990).
                                                                                                                                                                                Length 212;
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Pred. No. 2.7e-08;
---heq 1, Indels
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                                                                                          212 AA; 22831 MW; 948BADB72AAFA556 CRC64;
BY SIMILARITY.

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N-LINKED (GLCNAC. ..).
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94.7%; Pred. No.
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InterPro; IPR00169; SHprot acsite.
Pfan, PP00112; Peptidase_Cl; 1.
Pr0Dom, PD000158; PAPAIN.
ProDom, PD000158; Peptidase_Cl; 1.
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Best Local Similarity
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ARE DIFFERENT FROM THOSE OF OTHER CYSTEINS PROTEINASES.
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"Stem bromelain: amino acid sequence and implications for weak
minding of cystatin," 247,419-424 (1989).
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Rationja A., Rowan A.D., Buttle D.G., Rawlings N.D., Turk V.,
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Pred. No. 4.4e-10;
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     PROSITE, PS00640, THIOL PROTEASE ASN, 1.
Hydrolase, Thiol protease, Zymogen; Signal.
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                                                                                                                                      the Buropean Bioinformatics Institute. There are no restrictions on itse use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.iab-sib.ch/announce/or send an email to license@iab-sib.ch).
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Blochim. Bodybe, Acta 1345-231-236(199)
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15-JUN-2002 (Rel. 41, Last annotation update)
Cathepsin L precursor (EC 3.4.22.15).
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Ffam; PF00112; Peptidase_C1; 1.
PRINTS; PR00705; PAPAIN.
Probom; PD000158; Peptidase_C1; 1.
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PROSITE; PS00640, THIOL PROTEASE ASN;
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HSSP; O60911; 1FH0.
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                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: LYBOSOMAL.
SIMILARITY: BELONGS TO PEPTIDASE FAMILY CL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
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                                                                                           Query Match
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SIGNAL
PROPEP
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Ol-OCT-1993 (Rel. 27, Greated)
Ol-OCT-1993 (Rel. 27, Last sequence update)
Ol-OCT-1993 (Rel. 27, Last sequence update)
Cystelne proteinase II (EC 3.4.22.) (CCI Cystelne proteinase III (EC 3.4.22.) (CCI Carica candamarconsis
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                                                                                                                                         NON TER
                                                                                                                                                                                 TitesPro IPRO0668; Peptidase CI.
IntesPro IPRO0668; Peptidase CI.
IntesPro IPRO0122; Peptidase CI.
Pron PRO0112; Peptidase CI.
Pron PRO0125; Peptidase CI.
PRO0125; Peptidase CI.
PRO0125; PRO0125; PERIDASE NIS; PARTIAL
PRO
                                                                                                                                                                                                                                                                                                                                                       Biol
                                                                                                                                                                                                                                                                                                                                                                      Walrewens V., Jeziri M., van Beeumen J., Schnek A.G.,
Kleinschmidt T., Looze Y.,
Tsolation and preliminary characterization of the cysteine-
proceinases from the latex of Carica condamarcensis Hook.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, eudicotyledons; core eudicots; Rosidae,
eurosids II; Brassicales; Caricaceae; Carica.
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PROSITE; PS00639; THIOL PROTEASE HIS; 1.
PROSITE; PS00640; THIOL PROTEASE ASN; 1.
                                                                                                                             SECORNCE
                                                                                                                                                            ACT_SITE
                                                                                                                                                                                                                                                                                                MEROPS; C01.020;
                                                                                                                                                                                                                                                                                                               HSSP, PI40BO;

    -!- SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1

                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=94030669; PubMed=8216902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Latex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI Tax1D=29731;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P32956;
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                                                                                                                                                                                                                                                                                                                                           iol. Chem. Hoppe-Seyler 374:501-506(1993).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 VPQSIDWRDSGAVTSVKNQG 20
         PESIDWRKKGAVTPVKNQG 20
                                    POSIDWRDSGAVTSVKNQG 20
                                                                                                                             45 AA;
                                                                                                                                                         Thiol protease, Glycoprotein.
25 25 BY SIMILARITY
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17 POTENTIAL.
                                                                                                                             4636 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        37178 MW;
                                                                                  78.981
                                                                                               78.14;
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CATHERSIN LIGHT CHAIN.
BY SIMILARITY.
INTERCHAIN (BY SIMILARITY).
                                                                               Score 82; DB 1;
Pred. No. 3.2e-07;
                                                                                                                             F4C5D2881886E291 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 83; DB 1;
Pred. No. 2.1e-06;
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N-LINKED (GLCNAC. . .)
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CATHERSIN & HEAVY CHAIN.
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                                                                   Migmatches
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                                                                                               Length 43;
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This arks store for marry is copycidit. It is produced through a collaboration between the Swiss institute of Bioinformatics and the DBML outstation. The Burgeau Boniformatical institute. There are no restrictions on its modified and that a resemble about a some of the second of th
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State of the properly-dependage activity.
                                                                                                                                                                                                                                                                                                                            MEDIINE=6703368], PubMed=3533924;
STRIDY D.A., Estkano A.H., Kachan J., Ravetch J.V., Unkeless J.C.;
"Cloring and characterization of a mouse eysteine proteinsse.";
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PROSITE: PROSIDE: PROFISSE C76; 1.

PROSITE: PROSIDE: PROFISSE C76; 1.

PROSITE: PROSIDE: PROFISSE C76; 1.

PROSIDE: PROFISSE C76; 1.

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PROFISE: 
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Plerre R.S., Materbusse P. Nilson-Hamilton M.
Close relationship of the majore excreted protein of transformed
murine (blooplasts to thiol-dépendent cathépaisms.").
Zance Res et 46(1590-451) 1986.
Le Cancer Res et 46(1590-451) 1986.
Le Charles MADRANA FOR THE OVERALL DESADATION OF PROTEINS IN
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*Comparison of cathegein L synthesized by normal and transformed
Acell at the gene, message, proceful, and oligosaccharide levels.";
Acell, Biochem. Biophys. 283:447-457(1990).
murine preprocathepsin L. An abundant transcript induced by transformation of Enbohasans", J. Clin. Invest. 81:1621-1629(1988).
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InterPror, IRROGOG68; Peptidase Cl.
InterPro; IRROGOG69; SHyrot acsite.
Pfam; PF00112; Pegcidase_Cl7.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=91112761; PubMed=2275556;
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EMBL, J02583, AAA37445.1, ...
EMBL, M20495, AAA39984.1, ...
EMBL, X04392, CAA27980.1, ...
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PIR, S13890; S13890.
HSSP, P07711; 1CJL.
                                                                                                                                                                                                                                                          SEGUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bakaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermetophyta, Magnoliophyta, eudicoryledons, core eudicots, Rosidae;
eurosids II; Brasscales, Caricacaes, Carica
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11.0X4:1996 (Ret. 16, last sequence update)
CALF-UNR-2005 (Ret. 11, last sequence to Major excreted protein) (MEP),
CALF-UNR-2005 (Ret. 4, last campocation update)
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Joseph L.J., Chang L.C., Staenkovich D., Sukhatme V.P.,
"Complete nucleoxide and deduced amno acid sequences of human and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryotai Mesazoa; Chordata; Cramiata: Vertebrata; Buteleostomi;
Mammalia: Butheria: Rodentia: Schurognathi; Muridae; Musinae; Mus
NCBI_TRAID=10990;
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Troom B.R., Gall S., Gottesman M.M.
'Sequence and expression of the cDNA for MRP (major excreted
processin's a transformation-regulated secreted cathepain.")
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Pred. No. 3.2e-07;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                              01-0CT-1993 (Rel. 27, Created)
01-0CT-1993 (Rel. 27, Last sequence update)
15-UNV-202 (Rel. 41, Last annotation update)
Cysteine proteinase IV (RC 3-4.22.-) (CC-IV) (Fragment).
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InterPro, IPR000169, SHprot_acsite.
Pfam: PF00112; Peptidase_C1; 1.
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                                                                                                                                                                                                                                                                        STANDARD:
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SEITH A.M., DAITON, J.P., CHOMPH. K.A. Kilbane C.L., Harrop S.A.,
MEAL W. S. Mitchley P.J. S., CHOMPH. K.A. Kilbane C.L., Harrop S.A.,
WEAL W. S. Mitchley P.J. S. 
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Eukaryota, Metazoa, Platyhelminth
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PROSITE; PS00139; THIOL PROTEASE TYS; 1.
PROSITE; PS00639; THIOL PROTEASE HIS; 1.
PROSITE; PS00640; THIOL PROTEASE ASN; 1.
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SUBCELLULAR LOCATION: Lysosomal.
SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1.
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(Rel. 41, Last annotation updat
precursor (EC 3.4.22.15) (SMCL)
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9CFE68A3F1193479 CRC64;
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MEDIALE-2008387, PANEND-C. D. Chem. T.P., Bantico M.-I., Toyn C.D.
MEDIA. (Aug. 18., Rounal by S.D.) Chem. T.P., Bantico M.-I., Toyn C.D.
Benil C.F., Ketchen K.A. Lee, J.J., Sonning C.M., Koo H.L.,
Benil C.F., Ketchen K.A. Lee, J.J., Sonning C.M., Koo H.L.,
McSfex K.S., Coomin L.A., Sher M., Pai G., Van Aken S., Daywa L.,
Tallon L.J., Gill J.K., Adams M.O., Cerren A.G., Drayes D.,
Goodman H.M., Somerville C.S., Copenhawer G.P., Press D.,
McEman M.C., White O., Elsen J.A., Shibburg S.D., Fress D.,
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The European Profit Transitution as floop as a second set it is so way modified and this settlement as more than a content of the content of t
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This style STORY metry is copyright. It is produced theory a collideration between the State intelligence of Bioinformatics and the Botto outcetton. The Baropean boinformatical intuitive. There as no restrictions on its modified and Dail's resentitions do not seen that the Thomas are content at it not only medited and Dail's resentitions enough a base content at it not only entities a price and agreement (see http://www.isb-sib.ci/cannounce/or end an equal to licenseedsb-sib.ch).
(POTENTIAL).
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30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cysteine proteinase (P-II (EC 3.4.22.-).
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EMBL, AL161594; CAB80572.1; --
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MEROPS; COL.022; --
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"Strupture and expression of two genee that encode distinct drought-
inducible cyseline proteinases in Arabidopsis thaliana.";
@mm 129:175-128(1993).
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                     ACTIVATION DEPTINE (DOTENTIAL)
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70.0%; Pred. No. 4.8e-06;
ive 2; Mismatches 4; Indels
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RUISA OR AFRGS909 OR F19422.190.
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J. Biol. Chem. 265116897-18997(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-1992 (Rei. 22 Created)
01-MAY-1992 (Rei. 22 Last sequence update)
15-UTW-2002 (Rei. 42 Last semptation update)
01yaanin beta chain precursor (EC 3.4.22.-).
01yaa setitu (Rice)
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InterPro; IPR000169; SHprot_assite.
Pfam; PF00112; Peptidase_C1; 1.
PRUNTS; PR00705; PAPAIN.
PROSER; PR00138; Peptidase_C1; 1.
PROSER; P800139; THIOL_PROTEASE_CTS;
PROSER; P800139; THIOL_PROTEASE_CTS;
PROSER; P800139; THIOL_PROTEASE_CTS;
      This SWISS-PROT entry is copyright. It is produced through a collaboration
                                                             -i- TISSUE SPECIFICITY: EXPRESSED ONLY IN SEEDS
-i- INDUCTION: BY GIBBERELLIC ACID (GA).
-i- SIMILARITY: BELONGS TO PEPTIDASE FAMILY CI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Sukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnollophyta, Liliopsida, Foales, Poaceae; Ehrhartoidae, Oryzeae, Oryze.
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PROSITE; PS00640; THIOL_PROTENSE_ASN; FALSE_NEG
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    -!- CATALYTIC ACTIVITY: Preferential cleavage with a

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Sizur J. Siochem. Z67:12546-1256(2000).
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                                                             SIMILARITY BELONGS TO PEPFIDASE FAMILY CI. AS CAUTION: THE AUTHORS REGARD THE SEQUENCE AS TENTATIVE, AS BELIEVE THAT IT MAY HAVE BEEN CONTAMINATED BY A HOMOLOGOUS
                                    PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'APFEGENTER'
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InterPro; IPRO00118; Gramulin,
InterPro; IPRO00168; Peptidame C.1.
InterPro; IPRO00169; SHproc; acrise,
Pro; Peptidame C.1; 1.
Pram; PPO0139; gramulin; 1.
Prints; PPO0139; PAPLIN;
PRINTS; PPO0139; PAPLIN;
PRINTS; PPO0138; PEPTIDAME C.1; 1.
SAMRT; SHO0277; G.NN; 1.
SAMRT; SHO0277; G.NN; 1.
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or send an easil to licensesisb-sib.ch).
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use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by an
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This sties story entry in copyright. It is produced through a colliberation between the Suisa institute of Bioinformatics and the BBEL obtaining on the Barpage and poinformatics of Belliuse. For ser are no restrictions on its user by comparing institutions as long as its content, is in no way mentione entry and a suisability of the service of the se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TSTRAINSEQUENCE - TOUR STATE - 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "TISSNO EXPERITATION" BOTH BARNE ALEMENT I. A RE-
found in the upper epidernia. The lower spidernia predaminatly
contain by recedeball. In semisification tublice expression is
greater at Engels VI-VII than at Stages IX XIII.
- INVENTING Expression in Serial ice Supressed by gene cells.
- SINGITATIV: BELOSK'S TO PEPTINGS TANIV CI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SUBCELLULAR LOCATION: Procathepsin L is secreted. Mature cathepsin
                                                                                                                                                                                                                Boujtad N., Ogwoegbu S.O., Garmier M., Lee C.-H., Martin B.M.,
Pagestycholog v. a stimulator of steroid hormone synthesis isolated
"Agentichology".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of serondopenses by THVRE.

CTALATIC ACTIVITY. Specificity close to that of papash. As compared to cathogain a cathogae activity rowards protein substrates, but has little activity on 2-kay-kay-kay-make, and no peptidy-lappediase activity.

Whenc, and no peptidy-lappediase activity.
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RRGSITE, REGOLS, FITOL POTENSE C1: 1.
RRGSITE, REGOLS, FITOL POTENSE C1: 1.
RRGSITE, REGOLGS, FITOL POTENSE L1: 1.
RRGSITE, REGOLGS, TITOL POTENSE LNS: 1.
SIGNAL, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEBS (act, 236.57-611698).
-1-FUNCTION: Important for the overall degradation of proteins in TAMCTION: Proceeding the form of the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CATHERS I. HEAVY CHAIN.
CATHERS I. LIGHT CHAIN.
BY SIMILARITY.
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             SEQUENCE OF 18-37, FUNCTION, AND SUBCELLULAR LOCATION STRAINS-SPREAMS-SPREAMS-STRAINS-SEQUENCE. WHEN SERVING SECTION COLIS;
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InterPro; IPR000169; SBprot, acsite.
PEam; PF00112; Peptidase_Cl; 1.
PRINTS; PR00705; PAPAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-88296890; PubMed=3402618;
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EMBL, S85184, AAB21516.1, -.
PIR, S07098, KHRTL.
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Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus
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Efficken Lawrence M., Zabludoff S.D., Wright W.M.;
"Cyolic proteins, a secretory graduct of rat Serioli cells, is the
Decomyre force of cathegain I.",
Mol. Endocrinol. 5:1785-1798 (1991).
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"Molecular cloring and sequencing of cDNA for rat cathepsin L.";
FBBS Lett. 23:169-73(1987)
                                                                                                                                                                                                                                                                        PROSTITE, PROFESSOR, PRECISENCE, PALSE NRC.
PROSTITE, PROFESSOR, PROFESSOR, PALSE NRC.
PROSTITE, PROFESSOR, PROFESSOR, PALSE NRC.
PROSTITE, PROFESSOR, PRO
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MEDLINE-9005243; PubMed-2599113;
Nedne structure and 5' upstream sequence of rat cathepsin L.";
PEBS Lett. 259:71-74(1989).
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WEDLINE=88030047; PLMBG6=3666143;
MEDLINE=88030047; PLMBG6=3666143;
MEDIGON K., TOWECET T., Imajoh S., Kawesaki H., Kominami E.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 221;
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70.0%; Pred. No. 6.1e-06;
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InterPro; IPR000668, Peptidase_Cl.
InterPro; IPR000169; SHprot acsite.
Pfam; PF00112; Peptidase_Cl7_1.
PRANTS; PR00705; PAPAIN.
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MEDLINE=21255611; PubMed=11356678;
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TISSUE=Sertoli cells;
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Best Local Similarity
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MEDLINE-90269230; PubMed=2347312;
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Turk V.;
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"The complete amino acid sequence of papa
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Bukaryota, Viridajlanias, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidae;
euroaida 11, Emsasicales; Caridaceae; Carida
Yamamoto B., Tomoo K., Doi M., Ohishi H., Yamamoto D., Tsuboi S., Okamoto H., Okada
                                                    MEDLINE=93075728; FubMed=1445868;
                                                                                                                                    "The refined 2.4 A X-ray crystal structure of recombinant human stefin B in complax with the cysteine proteinase papain; a nove of proteinase hibbitor interaction.";
EMBO J. 9:1939-1947(1990).
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15-JUN-2002 (Rel. 41, Last annotation update)
Papain precursor (EC 3.4.22.2) (Papaya proteinase
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NGDLIMES 9314560; Paulocaski K.;

NGOLIMEN M.; Nompachi Schoozski K.;

STANCHUR and expression of two genes that encode distinct drought-inducible greated proteinsees in Arabidopeis thallans.";

Gene 199:175-18211991
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10-807-1995 [Rel. 32, Last requence update)

15-078-2002 [Rel. 41, Last annotation update)

Cystein proteinse RD13A precursor [RC 3.4.22.-),

RD13A OR AVKG3993 OR F19122.190.
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L -> I (IN REF. 2).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         123 VPOSIDWRDSGAVTSVKNO 141
                                                                                                                                                                                                                                                                                                                                                                                                                                      1 VPQSIDWRDYGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD:
                                                                                                                                                                        345 AA,
                                                                                                                                                                                                                                                                                                       Similarity
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        178
273
291
324
                                                                                                                                                                                                                                                                                                                                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RD19 ARATH
ID RD19 ARATH
AC P43296;
        DISULPID
                                                                                            CONFI, TOT
                                                                                                                                                                                 SECUENCE
                                                                                                                                       CONFLICT
                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                 datches
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        REFER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota, Viridipliantes, Streptophyta, Embryophyta, Tracheophyta,
Spermacophyta, Magnoliophyta, Liliopsida, Commelinidae incertae sedis,
Bromaliaceae, Ananas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Competide bands Best reported sessi materials ubstrate Bz-Phr. Val. Arg. | Weben but bands specified session materials and arguerate by the session se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CATALYTIC ACTIVITY: Hydrolysis of proteins with broad specificity
                                                                                            Gana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-9745461; Pubhwd-9315753; 

MER K.L., Albert K.L., Bernacocht R.J., Edmunds T.; 

"Complete amino acid sequence of annuin and a comparison with stem 

"Complete amino acid sequence of annuin and a comparison with stem 

"Depositati and "Other plant or operation".
                                                                                    ..
        100.0%; Score 108; DB 1; Length 212;
100.0%; Pred. No. 2.5e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=cv. Smooth Cayenne, TISSUE-Stem,
Robertson C. E., Goodenough P. M.,
"Cloning and expression of ananain gene from pineapple.";
Submitted (NOV-1997) to the BMBL/Genhank/DDBJ databases.
                                                                                    0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          - TISSUE SPECIFICITY: STEM
- MASS SPECIFORETRY: MAZ3478, METHOD=Electrospray.
- SIMILARITY: BELONGS TO PRETIDASE FAMILY CI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1997 [Rel. 35, Created]
L6-OCT-2001 [Rel. 40, Last sequence update)
L5-JUN-2002 [Rel. 41, Last annotation update]
Annahan precursor [RC 3.4.22.31)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT; 345 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BY SIMILARITY.
BY SIMILARITY.
BY SIMILARITY.
                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      POTENTIAL.
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InterPro, IPR000169; SHprot aceite.
Pfam; PF00112; Peptidase Cl. 1.
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                                                                                                                                                                        1 VPOSIDWRDYGAVTSVKNON 20
                                                                                                                                                                                                                                   2 VPOSTDMRDYGAVTSVKNON 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ananas comosus (Pineapple)
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                                                                                            20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 123-338.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P14080: 1YAL.
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PB0884; 022293;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Stem
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ACT SITE
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DISULFID
CARBOHYD
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Latralla B., Courtwy I., Cloud J., Abbotc A., Bootc K., Onharon D.,
Latralla B., Courtwy I., Cloud J., Abbotc A., Bootc K.,
Kidner J., Bablay D., Patido B., Miares K., Gegin K., Hiller L.,
Malton J., Spatch J., Wan E., Andrews S., Gesta G., Layana D.,
Malton J., Spatch J., Wan E., Maltowes S., Costa G., Layana D.,
Malton J., Spatch R., Johnson K., Doron K., Octrom M., Goshu C.,
Antonou B., Zidner B., Spatch J., Gashi H., Johnson D.,
Malton J., Bablay B., Spatch J., Malton J., Malton J.,
Sachy I.K., O'Shandhhusy A., Rodrigner M., Hoffman J., Till S.,
Genan S., Sholdy N., Massgaan A., Meccanic W. R.,
Marca M., Marca M., Massgaan A., Maccanic W. R.,
Marca M., Marca M., Marca M., Maccanic W. R.,
Marca M., Marca M., Marca M., Marca M., Johnson A., Ondon S., Marca M., Marc
                                          0.1400/1397 (8a.). 35, Leat meguance update)
11.300/2002 (8a.) 41, Leat monotarion update)
12.300/2002 (8a.) 41, Leat monotarion update)
0.1100/2002 (8a.) 41, Leat monotarion update (8a.) 41, Leat monotarion update)
41, Leat monotarion update (8a.) 41, Leat monotarion update)
41, Leat monotarion update (8a.) 41, Leat monotarion update)
41, Leat monotarion update (8a.) 41, Leat monotarion update)
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                                                                                                                                                                                                                                                                                                                                                            01-JAN-1990
01-NOV-1997
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PROSITE, PRO0138; PEDITIONS C1; 1.

PROSITE; PRO0139; THIOL PROTENSE CYS; 1.

PROSITE; PRO0639; THIOL_PROTENSE_HS; 1.

PROSITE; PRO0640; THIOL_PROTENSE_NSN; 1.
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InterPro; IPRO00169; SHprot_acsite.
Pfam; PF00112; Peptidase_Cl; 1.
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          NCBI_TaxID=3649;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 VPQSIDWRDYGAVTSVKNQ 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CARPA
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                                                                                                                                                                                                                                                                                                                 (Rel. 13, Created)
(Rel. 35, Last seq
(Rel. 41, Last ann
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329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72.24;
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CYSTENE PROFESINGE RD19A.

RY SIMILARITY.

BY 
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Pred. No. 3.2e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    352 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 368;
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          CATL_PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  무
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          CATL PIG
Q28944,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDINE-9712865; Pabhed-997205; Whyna L., Looze Y.;
Wess D., Bouckest D. Strip Cremes F. Wyna L., Looze Y.;
Berneture of chypopalation of the control of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Watson D.C., Yaguchi M., Lynn K.R.,
"The amino ecid sequence of chymopapain from Carica papaya.",
Biochem, J. 266:75-81(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hydrolame; Thiol protesse; Zymogan; Signal; 3D-structure.
SIGNAL 1 1880 ZYMOGENTIAL
PROPER 19 134 ACTIVATION PEPTIDE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TinterFro, IRRO0159; SHorot aceite.
Fram: PRO0112; Peptidase C11 1.
PROWNERS PRO0105; PAPAIN
PROMES: PRO0105; Peptidase C1; 1.
PROSITE; PRO0139; THIOL PROTEASE HIS; 1.
PROSITE; PRO0639; THIOL PROTEASE HIS; 1.
PROSITE; PRO0639; THIOL PROTEASE ANS; 1.
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PIR, S08285; S08285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; X97789; CAA66378.1; -.
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Biol. Chem. Hoppe-Seyler 370:425-434(1989)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jacquet A., Kleinschmidt T., Schnek A.G., Looze Y., Braunitzer G.:
"The thiol proteinases from the latex of Carica papaya L. III. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=89302685; PubMed=2500950;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90179730;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

    -!- CATALYTIC ACTIVITY: Specificity similar to that of papain.
    -!- SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1.

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                                                                                                                                                                                                                                                                                                                            136 POSIDWRAKGAVTPVKNQ 153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CONSELACINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             352 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    70.4%; Score 76; DB 1; Length 35:
83.3%; Pred. No. 6.4e-05;
Live 0; Mismatches 3; Indels
                                                                         PRT
                                                                         339 AA
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SECUENCE FROM N.A. Sus scrofa (Pig) NCBI TaxID=9823;

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Bukaryota, Viridiplantae, Streptophyta, Bmbryophyta, Tracheophyta,
Spermatophyta, Magnollophyta, eudicotyjedoms, core eudicote, Rosidae,
euroside II, Brassicales, Caricaçee, Carica,
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Spermatophyra, Magnoliophyra, eudicotyledons, core eudicots, Rosidae,
eurosida II; Brassloales, Carlocese, Carlos.
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                                                                                                                   Maltonevers V., Jaziti M., van Beenenn U., Schunk A.G., Maltonevers V., Jaziti M., van Beenenn U., Schunk A.G., Maltonevers V., Jaziti M., van Beenenn U., Schunk A.G., Standbricher M., Cooce V., Looce V., Looce V., Looce V., Cooce V., C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-027-1993 (Rel. 27) Last sequence update)
10-027-1993 (Rel. 27) Last sequence update)
15-UNN-2002 (Rel. 41, Last semotation update)
Cartica candamarcensis [11 (EC 3.4.22.*) (GC-III) (Fragment)
Cartica candamarcensis.
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Pred. No. 1.5e-05;
1; Mismatches 3; Indels
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-!- SIMILARITY: BELONGS TO PEPTIDASE PAMILY C1
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PROSTIE: PS00649; PHIOL PROTEASE AS; PARTIAL PROSTIE: PS00640; THIOL PROTEASE AS; PARTIAL PROSITE: PS00139; THIOL PROTEASE CYS; 1.
                                                                                                                                                                                                                                  43 AA
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                                                                                                                                                                                                              PRT;
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InterPro, IPR000169, SHprot acsite.
                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-1993 (Rel. 27, Created)
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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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- PROMITTINE INFORMAT FOR THE OVERALL DEGRADATION OF MOTERIAS IN
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                                                                                                                                                                                                          Stearns N.A., Dong J., Pan J.X., Brenner D.A., Sahagian G.G., 
"Comparison of cathepisin is synthesized by normal and transformed 
calls at the gene, message, protein, and oligomaccharide levels.", 
Arch. Biochem. Biophys. 23:447-457(1990).
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TISSUE=Liver;
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REDLINE=87033683; PubMed=3533924;

ROTINOP D.A., Erickson A.H., Kochan J., Ravetch J.V., Unkeless J.C.;

Politing and characterization of a mouse cysteine proceinase.*;

Total Characterization of a mouse cysteine proceinase.*;
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"Complete nuclectide and deduced amino acid sequences of 
murine preprocathepain L. An abundant transcript induced 
transformation of fibroblasts.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Sequence and expression of the cDNA for MEP (major excreted protein), a transformation-regulated secreted cathepsin."; Biochem. J. 246:731-735(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
MEDLINE-88076849; PubMed=3689328;
Troen B.R., Gal S., Gottesman M.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse)
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                                                                                                                                                       STRAIN-BNL
                                                                                                                                                                       SEQUENCE OF 89-300 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                 J. BIOI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=88213715; PubMed=2835398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TAXID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             43 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                 261:14697-14703(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4685 MW;
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Last annotation update)
(EC 3.4.22.15) (Major excreted protein) (MEP)
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Pred. No. 1.5e-05
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-CAYALYTIC ACTIVITY: Specificity close to that of papain. As compared to osthopaus B. osthopaus B. osthobitis bigher activity coards protein substrates, but has little activity on 7-Arg-Arg-MeMer. and no peptidy-dipeptidase activity.
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This SHES FROT marry is copyright. It is produced through a collaboration between the bases instituted is additionantical and the BMD, outsettion. The BUDGER HES SHEET AND A CONTROL OF THE SHEET AND A SHEET AND -!- SUBCELLULAR LOCATION: Lysosomal.
-!- SIMILARITY: BELONGS TO PEPTIDASE FAMILY CL a collaboration -

EMBL; X06086; CAA29470.1; -.
EMBL; X02583; AAA37445.1; -.
EMBL; M20495; AAA3984.1; -.
EMBL; X04392; CAA27980.1; -.

MHRODES COL. 0.27 ...

MED DESTRUCTION OF THE PROPERTY OF THE

Thiol protease; Glycoprotein; Lysosome; Zymogen; Signal
1 17 POTENTIAL. ACTIVATION PERTIDE

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BY SINILARITY Pred. No. 0.00013 INTERCHAIN (BY SIMILARITY).
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M -> I (IN REF. 2).
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P43295; Q96UT5; 01-NOV-1995 (Rel. 32, Created) 16-OCT-2001 (Rel. 40, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) PKT? JOL AA

Probable cysteine proteinase A494 precursor (EC 3.4.22.-) AT2G21430 OR P3K23.19

Arabidopsis thaliam (Mouse-ear crees). Bikaryota, Viridajalanae, Etrepschitya, Embryophyta: Tracheophyta; Spermatophyta: Magnoliophyta, sudicotyledons; core eudicots; Rosidae, enrosida [1] Emsaticales; Brassicaces, Farbidopsis.

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115-288-1996 (Rel. 33, Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bukaryota, Metazoa, Platyhelminthes, Trematoda, Digenea, Strigeidida,
Schistosomatoidea, Schistosomatidae, Schistosoma.
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cathepsin L precursor (EC 3.4,22.15) (SMCL1).
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InterPro; IRR000169; SHprot acsite.
Edm; PF00112; Peptidase_Cl; 1.
PRINTS; PR00705; PAPAIN.
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                                                                     SEQUENCE FROM N. A. STATE FROM T. P. BERLED M. 11, TOWN C.D., PERSEAN REV. CALLED M. 11, TOWN C.D., PERSEAN S.P. DEBERLED M. 11, TOWN C.D., PERSEAN S.P. DEBERLED M. 11, TOWN C.D., SETTING C.D., SETTING C.D., SETTING C.D., PERSEAN S.P. TOWN C.D., SETTING C.D., PERSEAN S.P. TOWN C.D., SETTING C.D., PERSEAN S.P. TOWN C.D., SETTING C.D., SETTING C.D., SETTING C.D., CORRESON M. D., CERTERS D. M. SETTING C.D., SETTING C.D., SETTING C.D., SETTING D. S.D., SETTING D. S
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39819 MW; D09A2CE5654642DD CRC64;
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MEDLINE=94289649; PubMed=8018874;
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Bukartotakos; Oryzain Or
This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the BMEJ outsation the Baropean Bioinformatics Institute, on there are no restrictions on its the Baropean Bioinformatics Institutions, There are no restrictions on its use at its content, as in in way made by non-profit institutions as long as its content, as in in way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                STRAIN-O, Nipponbaro, TissUB-Soudi Media Media
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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PROSITE; PS00639; THIOL PROTEASE HIS;
PROSITE; PS00640; THIOL PROTEASE ASN
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Echinostomida, Echinostomata; Fascioloidea; Fasciolidae; Fasciola
NCBI_TaxID=6192;
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                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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                                                                                                                                        SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO PEPTIDASE FAMILY CI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Lysosomal (Potential).
DEVELOPMENTAL STAGE: EXPRESSED AT THE NEWLY EXCYSTED JUVENILI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBUNIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION:
                                                                                                                                                              INDUCTION: BY GIBBERELLIC ACID (GA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BONDS (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NHMec, and no peptidyl-dipeptidase activity.
SUBUNIT: DINER OF AN HEAVY AND A LIGHT CHAIN LINKED BY DISULFIDE
                                                                                                                                                                                                                                                                                                                                                                                       TaxID=4530;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VPASIDWREYGYVTEVKD 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OKYSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VPQSIDWRDYGAVTSVKN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biophys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 AA; 2242 NW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ., Ashman K., Meeusen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               THIOL PROTEASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            72.24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Commun. 213:169-174(1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 72;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              53FFB5835EBCB0D7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   471 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19,
                                                                                                                                                                                                                                                       of multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01_0CT_1993 (Rel. 27, Created)
01_0CT_1993 (Rel. 27, Last sequence update)
15_UN_2002 (Rel. 41, Last annotation update)
Cysteine proteinase II (EC 3.4.22.-) (CC-II) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC2
    Pfdam, PF00112; Peptidase C1; 1.
ProDom, PF000158; Pptidase C1; 1.
PROSITE: P500639; THIOL PROTEASE HIS:
PROSITE: P500640; THIOL PROTEASE ASN;
PROSITE: P500139; THIOL_PROTEASE CYS;
                                                                                                                                                                                                      proteinames from the latex of Carica candamarcensis Hook,
Biol. Chem. Hoppe-Swyler 174(501-506(193)).
11 SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1.
                                                                                                                                                                                                                                                       Walreavens V., Jaziri M., van Beeumen J., Schnek A.U.,
Kleinschmidt T., Looze Y.;
"Isolation and preliminary characterization of the cysteine-
"Isolation from the larey of Carica candamarcensis Hook,";
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bukaryota, Viridiplantae, Stroptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, eudiorbyladona;
eurosids II; Brassicales; Caricaceae; Carica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P32955;
                                                                                                                                                              MEROPS; COL
                                                                                                                                                                                                                                                                                                                                               MEDITNE=34030889; MUDMed=8218902;
                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=29731;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carica candamarcensis.
                                                                                                                  InterPro: IPR000668; Peptidase_Cl.
InterPro: IPR000169; SHprot_acsite.
                                                                                                                                                                                                                                                                                                                                                                     TISSUE-Latex;
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CARCN

STANDARD;

PRT;

43

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P14080; 1YAL.

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Matches
                                                                                Query Match
                                                                                                                                                                     DISULFID
                                                                                                               SECUENCE
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                                                                                                                                                       DISULFID
                                                                                                                                                                                                                                             CHAIN
                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00112; Peptidase_Cl
Pfam; PF00396; granulin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEROPS; C01.029.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSP; P00785;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                     Hydrolase;
                                                                                                                                                                                                                                                                                                                                                         ProDom, PD000158, Peptidase_C1, 1
                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000169; SHprot_acsite
                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro, IPRO00118;
                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR000668; Peptidase_Cl
140 LPESVDWREKGAVAPVKNQ 158
                                                                    Local
                         1 VPQSIDWRDYGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           D90407;
                                                                                                                                                                                                                                                                                                                                                 SM00277; GRAN;
                                                                                                                                                                                                                                                                                                                                                                             PRO0705; PAPAIN
                                                                    Similarity
                                                                                                                                                                                                                                                                                                      PS00640; THIOL PROTEASE ASN,
PS00139; THIOL PROTEASE CYS;
PS00639; THIOL PROTEASE HIS;
                                                                                                            161
195
295
340
386
471
                                                                                                                                                                                                                                                                                       TOTAL
                                                      Conservative
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                                                                                                                                                                                                                                                                                       procease;
                                                                 66.7%;
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BY SIMILARITY
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                                                                                                                                                                                                  BY SIMILARITY.
BY SIMILARITY.
                                                                                  Score
                                                                                                            N-LINKED (GLCNAC. . .) ()
N-LINKED (GLCNAC. . .) ()
N-BF39D33995CEBID CRC64
                                                                                                                                                                                                                                             ORYZAIN BETA CHAIN
                                                                                                                                                                                                                                                              ACTIVATION PEPTIDE (POTENTIAL)
                                                                                                                                                                                                                                                                         POTENTIAL
                                                         Mismatches
                                                                                  72;
                                                                       No. 0.00039;
                                                                                DB 1;
                                                                                  Length 471
                                                                                                                                         .) (PUTENTIAL)
                                                         Indels
                                                                                                                          (POTENTIAL)
                                                      Q.
                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida; Zingiberales; Zingiberaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DODG K.H. Laursen R.A.;
Postolic State of System and System of Cysteine protesses with proline specificity from ginger rhizome Zingiber officials."

1. - Stochem. 267:1595-1557(1000)

- CORANTIT ACTIVITY: Preferential clowage with a proline residue
                                                                                                                                                                                              Gabs
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                                                                                                                              th (Similarity 68.4%; Score 71; DB 1; Length 43; Statianity 68.4%; Pred. No. 4.7e-05; 13; Conservative 2; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        909A312BD8632D42 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | Glycon | Inc. 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-LINKED (GLCNAC. ..).
/FTIG=CAR 000190.
N-LINKED [GLCNAC. ..).
                                                                               1777ACB74BC5F34B CRC64:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        - :- SIMILARITY: BELONGS TO PEPTIDASE FAMILY CL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /FTId=CAR 000200.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 221 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT: 221 AA
                               BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HSSP, P00785, ZACT.
GlyCoSulteDB, P84744,
InterPro, IPR000689, Peptidase_Cl.
InterPro, JPR000169, Stproc acsite.
Pfam, PF00112, Peptidase_Cl. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=20156257; PubMed=10691991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              221 AA: 23922 MN:
                                                                               4568 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.78:
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                                                                                                                                                                                                                                              2 POSIDWRDYGAVTSVKNON 20
                                                                                                                                                                                                                                                                                             2 PGSVDWROKGAVTPVKDON 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 VPQSIDWRDYGAVTSVKNQ 19
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Hydrolase, Thiol protease.
ACT SITE 25 25
NON-TER 43 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       STANDARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                          43
43
43 AA;
                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI TaxID=94328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         156
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                                                                                                                                                                                                                                                                                                                                                                                                                                 GPII ZINOF
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P82473;
                                                                               SROTENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                         Marches
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Search completed: November 21, 2002, 16:09:01

Job time : 8.5 secs

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1 VPOSIDWRDYGAVTSVKNQ 19 3 LPDSIDWREKGAVVPVKNQ 21 Local Similarity 68.4 es 13; Conservative

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Query Match

Matches

Length 221;

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10-6MY-2000 (Rel.) 35, Lost sequence update)
15-0MY-2000 (Rel.) 45, Lost sequence update)
15-0MY-2000 (Rel.) 41, Lost sequence update upda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Choi K.H., Lauren R.N., Starten R.N., Spranger and glycan structures of cysteine proteases with proline specificity from of singer and singer and singer the Stronger of Siccinaie. "S Bur. J. Sicchem. 267:1319-1526 (2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -: - CATALYTIC ACTIVITY: Preferential cleavage with a proline residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-LINKED (GLCNAC. ..) (POTENTIAL)
N-LINKED (GLCNAC. ..) (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -i- SIMILARITY: BELONGS TO PEPTIDASE FAMILY C1.
-I- CAUTION: THE AUTHOROSE REGARD THE SEQUENCE AS TENTATIVE, AS T
-I- CAUTION: THE AUTHOROSE REGARD THE SEQUENCE AS TENTATIVE BY A HOMOLOGOUS
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65.7%; Pred. No. 0.00026;
Pred. No. 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ProDom, PD000158, Peptidase (1, 1)
PROSITE, PS001019, THIOL PROTEASE (YS) FALSE NBC, PROSITE, PS000139, THIOL PROTEASE HIS, FALSE NBC, PROSITE, PS00640; THIOL PROTEASE ASN, 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MACOLAGO TITLOLOGO TO THE CONTROL OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000669; Peptidase Cl.
InterPro; IPR001669; SHprot acsite.
Pfam, PP00112; Peptidase Cl. 1.
PRIMTS; PR00705; PABAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Root;
MEDLINE=20156257; PubMed=10691991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          221 AA; 24241 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HSSP; P00785; 2ACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zingiber.
NCBI TaxID-94328;
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length: 2000000000
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Q9S770;
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Q1-MAY-2000 (TYEMBLrel. 13, C)
Q1-MAY-2000 (TYEMBLrel. 13, L)
Q1-DEC-2001 (TYEMBLrel. 19, L)
BYONG-14IN (FYEMBER).
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Theories, Propositions C.T., Theories, Propositions C.T., Bernard C.T., Bernar
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MEDLINE=94330946; PubMed=8053898;
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MEDLINE=95298192; PubMed=7779262;
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Spermatophyta; Magnoliophyta; Lillopeida; Commelinidae incertae sedie;
Bromeliaceae; Ananas.
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                    1 VPQSIDWRDSGAVTSVKNQG
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Conservative 0;
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Bukaryota, Viridiplantās; Streptophyta; Embryophyta; Tracheophyta;
Spermocophyta; Magnaliophyta; Liliopsida; Commelinidae incertae sedis;
Escomilaceae; Ananas.
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Bikaryor, Vitidiplantes, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnollophyta, Bronzpophyta, Bromeliatese, Anana.
Bromeliacese, Anana.
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STRANDEV, NEV. D. SISSUE-FRUIT;
Mith E., Chamoto Y., Cha E.,
Coloning and sequencing of Cygecine proteinaee in Anamas comosus.";
Submitted (CGT-1994) to the SMGL/Genflank/DDB detabases.
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SIGNAL 1 36 CYPERINE PROTEINABE PRECURSOR, ANN SEQUENCE 356 AA, 35528 MM, 970705652EC01DBB CRC64;
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01-JAN-1998 (TYENBELE). 05, Last sequence update)
01-NAR-2002 (TYENBELE). 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Cysteine proteinase, ANS precursor.
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Pfem; PR00112, Peptidase_Cl; I.
PRNTS; PR00705, PAPAIN.
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Bukaryors, Viridijantas, Etreptophyta; Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, Lillopsida; Commelinidae incertae sedis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ananas comonguis (Rineappple).
Spermatophytes, Partiplaties; Streptophytes, Embryophytes, Tracheophytes;
Stormatophytes, Magnobitophytes, Lillopsida, Commellinidae incertae sedis;
Stormatophytes, Magnobitophytes, Lillopsida, Commellinidae incertae sedis;
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Maurer H.R.,
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"Bond.com" (Page 1971)
"InterPro Pool InterPro (Page 1971)
"The Pro Pool InterPro (Page 1971)
"The Pro (Page 197
                                                                                                                                                                                                                                                                   01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 17, Last amocration update)
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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01-NCV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last Sequence update)
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Matches 18; Conservative
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R PÉSIN, PRODOS; PANALITA 1.

R PESINS, PANALITA 1
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040922; 040919;
01-NOV-1996 (TERBLIZE). 01, Created)
01-NOV-1996 (TERBLIZE). 01, Last amnotation update)
01-MAR-2002 (TERBLIZE). 20, Last amnotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000118; Granulin.
InterPro; IPR000668; Paptidase_C1.
InterPro; IPR001211; PhospholipaseAZ.
InterPro; IPR00169; SHprot_acsite.
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EMBL; Z49765; CAA89835.1; -.
HSSP; P07711; ICJL.
MEROPS; C01.029; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE-MHOLE SEEDLING,
Tranbarger T.J., Misra S.;
Submitted (MAY-1995) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN- (MIRB) FRANCO;
Tranbarger T.J., Misra S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene 172:221-226(1995).
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STRAIN-(MIRR) FRANCH, STRAIN-
STRAIN-(MIRR) FRANCH-
TRADBETGET T.J., WRITE S.;

TRADBETGET T.J., WRITE S.;

SELECTION AND ADDRESSION Of a developmentally regulated cDNA encoding a cyptesting processes personce promotorabil from Douglas fir.";
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NCBI_TaxID=3357;
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Bukaryota, Viitdiplantae, Sureptophyta; Embryophyta; Tracheophyta
Spermatophyta, Coniferopsida; Coniferales; Pinaceae; Pseudotsuga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PM33CYSP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prystor. Plantarum 0:0-0(0).
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                                                                                                                                               132 LPESIDWREKGAVTAVKNOG 153
                                                                                                                                                                                                                                                      T VPQSIDWRDSGAVTSVKNQG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VPQSIDWRDSGAVTSVKNQ 19
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                                                                                                                                                                                                                                                                                                                                                   Conservative
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94.7%; Pred. No. 3.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                  Score 88; DB 10; Length 454;
Pred. No. 3.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FBSB
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Best Local Similarity
Query Macch
                                                                                                                                                                                                                       MEROES, COLIOCE:
InterFro, IPROCOSES, Paptidase_CL.
InterFro, IPROCOSES, SUpro_Accite.
IPROFFRO, IPROCOSES, SUpro_Accite.
IPROFFRO, IPROCOSES, SUpro_Accite.
IPROCOSES, PROCOSES, PERIOR SECTION FROM THOU FORTHASE CITS, INC.
IPROCITES, PROCOSES, THOU_ROTTHASE CITS, INC.
IPROCITES, PROCOSES, PROCESS, PROC
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Q9U554;
01-MAY-2000 (TEMBLES: 13,
01-MAY-2000 (TEMBLES: 13,
01-JUN-2002 (TEMBLES: 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE-LEAF; Al-Sheikh M., Revell D.F., Sumner I.G., Connerton I.F.;
Taylor M.A., Al-Sheikh M., Revell D.F., Sumner I.G., Connerton I.F.;
"CDNA cloning and expression of Carica papaya prochymopapain isoforms"
"CDNA cloning and expression of Carica papaya prochymopapain isoforms"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Carica papya (Papaya).
Bukayyota Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, eudicoryledons; core eudicots; Rosidae;
eurosids II; Brasscales; Caricaeses; Carica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             098089 PRELIMINAR, PRF, 226 AA.
098089 PRELIMINAR, PRF, 226 AA.
098089 01-04V-2000 (FRMELFal. 1) Located
01-04V-2000 (FRMELFal. 1) Located
01-04W-2000 (FRMELFAL 1) Located
01-04W-2000 (FRMELF
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1888) PO7711 1021.
1888) PO7711 1021.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tethya aurantia.
Eukaryota; Metazoa; Porifera; E
Hadromerida; Tethyidae; Tethya.
                                                                                                                                               Hydrolase; Thiol protease.
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EMBL; AJ131997; CAB38316.1; -.
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                                                                                            SEQUENCE
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                                                                                            226 AA; 24597 MW; 997A1B6E0CE49C35 CRC64;
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BU.UK; SCOTE 84;
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Ananae comeour (Pitaappia)
Ekkaryosa, Viridijlantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Spermatophyta; Liliopsida; Commellinidae incertae sedis;
Bromellinidae; Anana
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Application of the Computer of the Compu
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70.0%; Pred. No. 1e-05;
rive 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      80.0%; Score 84; DB 10; Length 324;
84.2%; Pred. No. 9.7e-06;
live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHAIN 95 312 FB1035.
SEQUENCE 324 AA, 36038 MM; 057462C59424339F CRC64.
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01-JUN-2002 (TERMBLeel. 21, Last sequence update)
01-JUN-2002 (TERMBLeel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                  Maddon (M.) Maddon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT: 337 AM
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"FORM - Cloting and expression of Carlos papaya prochymopapain isoforms
th Escherichia coll., "A. St. 44-47(1999).
"Blanc 601, 145-14-47(1999).
"RSSP, Artdeo, 17WL."
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"Cloring and sequencing of cysteine proteinases in Amanas comosus.";

"Cloring and sequencing of cysteine proteinases in Amanas comosus.";

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"BMBL: D35834; BAA22546.1;
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Best Local
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InterFor, IPRODOSS, Septidase_Cl.
InterFor, IPRODOSS, Septidase_Cl.
PRODOSS, Peptidase_Cl.
Prodos, PRODOSS, Peptidase_Cl.
Prodos, PRODOSS, Peptidase_Cl.
PROSITE, PRODOSS, PRODO
                                                                                                                 Matches
                                                                                                                                                                 Best Local
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01-MAY-2000 (TrEMSLrel. 13, Last sequence update)
01-MAR-2002 (TreMSLrel. 20, Last annotation update)
01-MAR-2002 (TreMSLrel. 20, Last annotation update)
01/MSDPpath isoform II precursor (SC 3.4.22.6).
CHMODIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9SMI1
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STRAIN-CV.MS-1.0, TISSUE-FRUIT,

WHILE E. Actanki H. Takata Y. Kono A., Okamoto Y., Oti-

Mind E. Actanki H. Takata Y. Kono A., Okamoto Y., Oti-

Cloning and expenseding of Eruit bromelain.*,

Submitted (JAN-1993) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                            PROBOTE: PRODOGLES: PRETIDATE CT: 1.

PROSITE: PRODESO: THIOL PROTEASE CTS: 1.

PROSITE: PRODESO: "HIOL PROTEASE CTS: 1.

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STORAL 26

POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR000668; Peptidase_C1.
InterPro; IPR000169; Shprot_acsite.
Pfam; PF00112; Peptidase_C1; 1.
PRINTS; PR00705; PAPAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TORMA cloning and expression of Carica papaya prochymopapain isoforms in Escherichia coli.";
Plant Sci. 145:41-47(1999).
EMBL, AJ131959; CABSB14-1J -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carica papaya (papaya).

Bukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicocyledoms; core eudicocus; Rosidae;
eurosids II; Brassicales; Caricaceae; Carica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSSP; P07711;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; D14059; BAA21849.1; -
                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEROPS; C01.002; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Taylor M.A., Al-Sheikh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=LEAF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=3649,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     123
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POSIDWRDSGAVTSVKNOG 20
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                                                                                                                         To:
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                                                                                                                                                                                                                                                                                   352 AA,
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                                                                                                                                                                                                                                                                                   39462 MW;
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                                                                                                                                                                 Pred. No.
                                                                                                                                                                                                                                                                                                                           CHYMOPAPAIN II
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                                                                                                                         Mismatches
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                                                                                                                                                                 1.1e-05;
                                                                                                                                                                                                  DB 10; Length 352;
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InterPro; PRODICES, Supro-golite.
Prodices PRODICES PROFILMS—CI.
PRODICES PROFILMS—CI.
PROFILMS—
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ol-MAY-2000 (TERMBLrel. 1, last exquence update)
Ol-MAY-2002 (TERMBLrel. 20, last amocation update)
Chymopapain isoform III precursor (EC 3.4.22.6).
CHYMOIII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q9SMI0;
Q9SMI0;
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1998 (TYEMBLrel. 08, Created)
01-NOV-1998 (TYEMBLrel. 08, Last sequence update)
01-MAR-2002 (TYEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carica papaya (Rapaya).
Bakaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta, eudicotyledom; core eudicots; Kosidae
eurosida II; Brassicales; Caricacese; Carica.
                                                                        "Cloning and expression of ananain gene from pineapple";
Submitted (JUT.1998) to the BMEL/GenBank/DDBJ databases.
BMBL, AJ009330; CAA08861.1; ".
BESSP, P14080; TAAL.
                                                                                                                                                                                                                                          STRAIN=CV. SMCOTH CAYENNE; TISSUE=STEM;
Robertson C.E., Goodenough P.W.;
                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=4615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnollophyta, Liliopsida, Commelinidae incertae se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anahas comosus (Pineappie)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cysteine proteinase, AN11 precursor.
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EMBL; AJ131996; CAB38315.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "cDNA cloning and expression
in Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Taylor M.A., Al-Sheikh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=LEAF;
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                                                MEROPS;
                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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Pred. No. 1.1e-05;
0; Mismatches 3
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Length 361; Indels 0

Gaps

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interPro;

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DE TILCATO-1 PROPOSES, PROFICIARE CIT.

PROPOSES, PROFICIA PROFICIARE CIT.

DE REGETTE, PROSES, PRICE, PROFILES, UNICARO, 1.

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Bananas conogas; (Pichogappale).
Speematoppaty Viridiplantes; Streptophyra; Embryophyra; Tracheophyra;
Speematophyra; Magnoliophyra; Liliopaidu, Cosmellinidae incertae sedia;
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100.0%; Pred. No. 1.88-09;
tive 0; Mismatches 0; Indels 0;
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90.0%; Pred. No. 5e-08;
ive 0; Mismatches 2; Indels
1 24 POTENTIAL.
123 338 FBSB.
356 AA; 39560 MW, 16C2DCCB47238928 CRC64;
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SEQUENCE 324 AA; 36038 MW; 057462C59424339F CRC64;
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01-JAN-1958 (TEBHBLEE) 05, Created
01-JAN-1958 (TEBHBLEE) 05, Last sequence update)
01-MAR-2002 (TEBHBLEE): 20, Last annotation update)
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01-JAN-1998 (TrEMBirel. 05, Last sequence update)
01-MAR-2002 (TrEMBirel. 00, Last annotation update)
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STRAIN=CV.N67-10; TISSUE=FRUIT;
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les 18, Conservative
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NCBI TaxID=4615;
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                                                                                                                                                                   Anamas cemceus (Pinaspile).

Makaryota, Yirdipjantae; Streptophyta; Embryophyta; Tsecheophyta; Spensicophyta; Magnallophyta; Liliopsida; Cosmellinidae incertae sedis; Respondatedor Anamas.

NGEL TRANDOAGES;
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Spermatophyra, Magnoliophyra, Liliopeida, Commelinidae incertae sedis;
Bromeliaceae, Ananas.
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WALE BY COMMENCE OF S. CHICATOR ESCALEDISE IN AGAINS CONDUMB-7;

SADMIETED GOT-2941 CER BENGLOGEN OF THE SERVICE OF SERVIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSTER; PROSECTO, THOLOUGH PRETAINS.
PROSTER; PROSTER, PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 108; DB 10; Length 356; 100.0%; Pred. No. 1.8e-09; 1.ve 0; Mismarches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-CY, SHOOTH CALENDE, TISSUE-STED,
STRAIN-CY, SHOOTH CALENDE, TISSUE-STED,
CALENTION CALENDER OF 
01-NOV-1998 (TrEMBLE). 08, Last sequence update)
01-MAR-2002 (TrEMBLE). 20, Last annotation update)
Cysteine proteinase, ANB precursor.
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01-JAN-1998 (TrEMBLrel. OS, Last sequence update)
01-JAN-1998 (TrEMBLrel. 20, Last annotation update)
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PROGRIEF PROGRIEG FRICTAGARMA, UNKNORM. 1.
PROGRIEF PROGRIEF PROGREGE AND 1.
PROGRIEF PROGREGE CYS; 1.
HATOL PROGREGE CYS; 1.
HATOL PROGREGE CYS; 1.
HATOL PROGREGE CYS; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERPROT IPRODOGES, PERLIDAGE CI.
INTERPROT, IPRODOLES, SHPRCL ACSIDE.
Pfam. PF00112, PEDLIDAGE CI. 1.
PRINTS, PRODOTOS, PAPAIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 VPOSIDWRDYGAVTSVKNON 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FBSB precursor.
Ananas comosus (Pineapple).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE PROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match
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Best Local S
Matches 18
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Best Local
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MERODES, COL.CORY...

MEROPO, MERODES, Poptidame CL.

MERODES, PRODULIS POPTIDAME CL.

PRINTERS, PRODUCES, POPALIN.

PRINTERS, PRODUCES, POPALIN.

PRODUCTION, PRODUCES, PARAMERA, PRODUCES, PRODU
                                    O. -MAY-2000 (TrBMBLrcd. 11) Late sequence update)
O1-MAY-2000 (TrBMBLrcd. 11) Late sequence update)
BComplain (TrBMBLrcd. 12) Late annotation update)
BComplain (Transparel)
Mananas comenus (Flesspple)
Subaryoca, Viridojalates (Streptophyta; Embryophyta
Spermatophyta; Magnolichyva; Liliopsida; Commelini
BCompliances; Annanas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Napper A.D., Bennett S.F., Borowski M., Boldridge M.B., Leonard M.J.,
Sogers E.S., Dekan Y., Laurenn R.A., Reinhold B., Shames S.L.;
Blochem. J. 301:727-752(1994).
InterFor, IPBODGES; Peptidase_Cl.;
Form. PPOD112; Peptidase_Cl.;
SEQUENCE 20 AJ, 2219 MW. 3384A16392CLEBC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OL-MAY-2000 (Trightize). 13. Created)
OL-MAY-2000 (Trightize). 13. Created)
OL-MAY-2000 (Trightize). 13. Las enteretion spidice)
OL-MAY-2000 (Trightize).
OL-MAY-2000 (Trig
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"Cloning and sequencing of fruit bromelain."
Submitted (JAN-1993) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE.
MEDLINE=94330946; PubMed=8053898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CARS6D
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NCBI_TaxID=4615;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPQSIDWRDYGAVTSVKNON 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         351 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91.7%; Score 99; DB 10; Length 35
90.0%; Pred. No. 5.4e-08;
Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 97; DB 10; Length 20
Pred. No. 4.6e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BROMELAIN
                                                                         eprophyta; zmbryophyta; tracheophyta;
billiopsida; Commelinidae incertae sedis;
                                                                                                                                                                                                                                                                                                                                                                 PKT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  POTENTIAL.
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                                                                                                              Tracheophyta
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                                                                                                                                                                                                                                                                                                                     Best Local
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INEAFTO, IPRODOGSS, POPILIANE CI.
INEAFTO, IPRODOSS, SHOPE, ARTHER
PERMITS, PRODOSS, PARAIN CI.
PECODOR, IPRODOSS, PARAIN CI.
PECODOR, PRODOSS, THIOW, PROTERSE_HIS, UNEXCON.
1.
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STRAINS-V., SECTAT CAYENNE, ITESUE-STEM;
STRAINS-V., SECTAT CONTROL OF THE PROPERTY AND STRAIN OF THE PROPERTY OF THE PROPERT
                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        081085;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=4615;
                                                                                                                                                   123 VPOSIDWRNYGAVTSVKN 140
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                                                                                                                                                                                                              1 VPQSIDWRDYGAVTSVKN 18
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                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                            27
27 AA;
                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                   39606 MW;
                                                                                                                                                                                                                                                                                                                     94.4%; Pred. No. 7.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CYSTEINE PROTEINASE PRECURSOR,
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POTENTIAL 0B3930D035E47C31 CRC64; Mismatches

Indels

0,

Gaps

0,

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Ananas comesus (Elmaepha).
Bakaryota; Wirdiplanias; Ersptophyta; Embryophyta; Tracheophyta;
Spermacophyta: Megnollophyta; hiliopsida; Commelinidae incertae sadis;
Bromeliace; Mananas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1998 (TERMBLTel. 08, Created)
01-NOV-1998 (TERMBLTel. 08, Last sequence update)
01-MAR-2002 (TERMBLTel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Napper A.D., Bennett S.P., Borowski M., Holdridge M.B.,
Rogers B.E., Duan Y., Laursen R.A., Reinhold B., Shames
Biochem, J. 301;727-735(1994).
InterPro, IPB00066B, Reptidase_C1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SECURNICE FROM M.A.
MEDIJER-95298122 Phaked=7779262;
MEDIJER-95298122 Phaked=7779262;
MENICHE H.R.;

                                                                                                                                                                                                                                                                                                                                                                                                                                    Cysteins proteinase, ANII precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ptam; PF00112; Peptidase_C1; 1.
SEQUENCE 20 AA; 2144 MW; 83A842BC812C3EBC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=94330946; PubMed=8053898;
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Pred. No. 2.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        357 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Inders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pisaciola gigantica.
Poblioperonidas Ecinoscomata; Fascioloidea; Prematoda; Digenea;
Poblioperonidas Ecinoscomata; Pascioloidea; Pasciolidae; Fasciola.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sobhon P., Meemon K., Grams R., Grams S.V., Korge G., Hofmann A., "Molecular cloning of cathepain L encoding genes from Fasciola of cathepain L.
                                                                                                                                                                                                                                                                                                                                                                               Length 352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74.1%; Score 80; DB 5; Length 326; 78.9%; Pred. No. 6.6e-05; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                        Query Match 83.3%; Score 90; DB 10; Length 352 Best Local Similarity 85.0%; Pred. No 1.6e-06; Indels Matches 17; Conservative 1; Masmatches 2; Indels Matches 17; Conservative 1; Masmatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   352 AA, 38823 MW; 5969AAC09EB3D367 CRC64,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        040822, 040819, 040818. 01, Created)
01-NOV-1996 (TREMBLEE), 01, Created)
01-NOV-2996 (TREMBLEE), 01, Laws Sequence update)
01-NAR-2002 (TREMBLEE), 02, Last annotation update)
BENDONTAIN (Cysteine processe).
PEGATE PRODUZE, PREPÉTIGASE CI: 1.
PRODES: PRODUSES, PERÉTIGASE CI: 1.
PROSTER. PRODUSES, PERÉTIGAS. 1.
PROSTER. PROSTESS, PITOL, PROTEASE CVS, 1.
PROSTER. PROSTESS, PITOL, PROTEASE CVS, 1.
PROSTER. PROSTESS, PITOL, PROTEASE FILS, DANSOHN 1.
PHOTOLOGIASE SIGNAL 1.
PATOL PROTEASE FILS, DANSOHN 1.
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                                                                                                                                                                                                                                                                              FR31 (PR13)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           124 VGOSIDWRDYGAVTEVKDON 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 VPQSIDWRDYGAVTSVKNQN 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VPOSIDWRDYGAVTSVKNQ 19
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Best Local Similarity 78.9%
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRELIMINARY:
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                                                                                                                                                                                                                                            340
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Q40922
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                                                                                                                                                                                                  Anana Compuse; (Pieneppple).
Semakepharyota, Viridiplatica; Szrepcophyra; Embryophyra; Tracheophyra;
Szemakephyra; Magnollophyta; Liliopaida; Commelinidae incertae sedia;
Bromeliacese; Atanas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ananas comesus (Finespopts).
Sermacophyta, Babryophyta, Tracheophyta; Sermacophyta; Barnacophyta, Manoliophyta, Jillopsida, Commellinidae incertae sedis; Bromediatese, Ananas.
                                                                                                                                                                                                                                                                                                                                                                                                                                propagation "W. (") The RESERVATE |

"Closing and sequencing of cytering proceinage in Anama concept.";

"Samitted (CGT-2594) to the BRA/CenSanK/DDM databases.

BRBH, DSSS11 BAAZS11 AAAZS14 | To the BRA/CenSanK/DDM databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
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MINTA E. AFARMAK! H. TARAEL W. KOND A., OKAMOLO Y., OTA S.,
SUUMILECA (AUG-1997) to the EMBL/Gonbank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               March 83.34; Score 90; DB 10; Length 340; Cocal Similarity 85.04; Pred. No. 1.6e-62; Indels 8 17; Conservative 1; Mismarches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
SERRAIN-CV. N67-10, TISSUE=FRUIT;
Mura E., Okamoco Y. Ota S.;
Submitted (OCT-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            340 AA: 37646 MW: 005586B006AA903F CRC64;
                                  021800;
01.-7AN-1998 (TERMELRE), 05, Created)
01.-7AN-1998 (TERMELRE), 05, Last sequence update)
01.-MAR-2002 (TERMELRE), 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JAN-1998 (TrEMBirel. 05, Created)
01-JAN-1998 (TrEMBirel. 05, Last sequence update)
01-MAR-2002 (TrEMBirel. 20, Last senotation update)
RB31 precursor (FB13 precursor) (Bromelain).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT: 352 AA.
      PRT: 340 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    123 VGOSIDWRDYGAVTEVKDON 142
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EMBL, D14058, BAA21848.1;
EMBL, D14057, BAA21929.1;
HSSP: P14080; IYAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 VPQSIDWRDYGAVTSVKNQN 20
                                                                                                                                                                      FB22 precursor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       >340
      PRELIMINARY:
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                                                                                                                                                                                                                                                                                                                                      NCBI TaxID=4615;
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SEOUENCE
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      023800
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MATURE IMAGINAL DISKS DEVELOPMENT

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SOR REPRESENTATION OF THE PROPERTY OF THE PROP
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y SECURED FROM N.A., PARTIAL SECURED. AND CHARACTERIZATION.

X MEDIAN K.F.I., MOTES S., MACCY S. I.

** HORMAN K.F.I., MOTES S., MACCY S. I.

** Exception for the secure of 
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Best Local Similarity
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01. Ward and (Tribulare). 01. Last sequence update)
01. Ward and (Tribulare). 01. Last sequence update)
02. Republic L presenter (ES 1.4.2.55).
03. Recording Recepting (Tribulary). (Restablished personnel).
04. Recording Recepting (Tribulary).
05. Recording Recording (Tribulary).
06. Recording (Recording Recording Rec
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Q26636,
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MEDILES-9625408; Dubwel-662107;
Franbarger T.J., Miera S.;
"Ernburte and expression of a developmentally regulated cDMA encoding of CPMA in processor of a developmentally regulated cDMA encoding a Cystaine processe (pseudotzain) from Douglas fir.";
Gene 172:221-261 (1995).
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PRODOS, PRODOS, PAPAIN.

PRODOS, PRODOS, PAPAIN.

PROSITE PRODOS, PAPAIN.

PROSITE PRODOS, PHOLOPESE ASN.

PROSITE PRODOS, PHOLOPESE TO PROSITE PRODOS, PROSITE PRODOS, PHOLOPESE TO PROSITE PRODOS, PROD
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Physiol. Plantarum 0:0-0(0).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00396; granulin; 1.
Pfam; PF00112; Peptidase_C1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUB-WHOLE SEEDLING,
Tranbarger T.J., Misra S.;
Submitted (MAY-1995) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           132
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InterPro, IPR001211; PhospholipaseA2
InterPro, IPR000169; SHprot_acsite.
Pfam: PF00396. granulin. 1
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Pred. No. 9.7e-05;
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Spermatophyta, Asgnoliophyta, endotyladoma, cudiocta; Rosidae,
Tracheophyta, Parasicales; Brassicaceae, Arabidopsis.
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Pfam, PP00112; Pepridase_C1; 1.
PRINTS; PR00105; PAPAIN.
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Artiguenave P., Saurin N., Neissenbach J., Newes H.W., Ruc
Lomcke K., Mayer K.F.X., Quetier F., Salanoubat M.;
Submitted (APR-1999) to the EMBL/GenBank/DDBJ databases.
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DEVELOPMENTAL STRAGE: HIGHLY EXPRESSED DURING EMERCONIC DEVELOPMENTAL STRAGE: HIGHLY EXPRESSED DURING EMERCONIC DEVELOPMENTAL STRAGE HIGHLY EXPENSES TO REPUTIONSE FAMILY C1; ALSO KNOWN AS THE DAMAIN FAMILY OF THIOL PROTESSES.
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SUBCELLULAR LOCATION: PROCATHEPSIN L IS SECRETED BY WHEN CULTURED IN THE PRESENCE OF 2-HYDROXYECDYSONE.
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THEREPOT INTRODUCES, PROTECT AND THE PROTECT A
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SEQUENCE 311 AA, 35409 MW; BB38A5DCB0510845 CRC64;
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01.NAR-2001 (TrEMBLrel. 16, Created)
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mail (EC ).4.22.31) procure mail (EC ).4.22.31) procure goods [pink] pro	Overy Macch malarity 100.01; Score 105; DB 2; Length 20; Macches 100 conservative 100; DB 2; Length 20; Macches 100; Macches 10; Indels 0; Gaps 0; Oy 1 POSITHOSOAVISTANCE 20  Db 1 POSITHOSOAVISTANCE 20	Bablin 1  Saddon (C. ).4.22.31) - pineapple (fragment)  Sandan (GC.).4.22.31) - pineapple (fragment)  C. Species Annua comesus (pineapple)  C. Species (Frb.).28; Seequardevision (1-Nov-1956 Htaxchange (7-May-1959)  C. Date: 1.6.78b.).28; Seequardevision (1-Nov-1956 Htaxchange (7-May-1959)  F. Napper. J. D. Sendat. S. S. S.; Becoment. N.; Holdridge, N. B.; Leonard. N.J.C.; Sequerial (1-Nov-1958)  F. Napper. J. Date: 1.5. Sept. Security (1-Nov-1958)  A. Accession (960) 8600; NOID 931696; PMIL865898  A. Molecule (1-Proper protein A. Maschdose) (1-O OAP)  A. Molecule (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-O OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-OAP)  A. Sept. Security (1-Proper protein A. Maschdose) (1-OAP)	, a		

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C;Accession: T10503
R;Muta. B.; Aramaki, H.; Takata, Y.; Kono, A.; Okamoto, Y.; Ota, submitted to the EMBL Data Library, January 1993
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C;Bate: 16-U1-1999 #sequence_revision 16-U1-1999 #text_change 20-Unn-2000
C;Accession: 7U500)
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A;Molecule type: mRNA
A;Residues: 1-224 «MIT»
A;Cross-references: EMBL:D38534
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submitted to the EMHL Data Library, October 1934
A. Description: Cloning and sequencing of cysteine proteinases in Amanas comosus
A. Reference number: 2.17065
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8;20-33/Domain: carboxyl-terminal propertion status predicted <CTP
8;30-33/Domain: carboxyl-terminal propertion status predicted <CTP
8;30-33/Domain: carboxyl-terminal propertion status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Cross-references: SMBL:D14059
A; Experimental source: cv. N67-10; nearly mature fruit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; MOIECUIE Type: mRNA
A; Residues: 1-351 < MUT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Description: Cloning and sequencing of fruit bromelain A; Reference number: Z17060 A; Accession: T10503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fruit bromelain (EC 3.4.22.33) FB18 precursor - pineapple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T10503
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eruit bromelain (EC 3.4.22.33) FB1035 precursor - pineapple (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Description: hydrolyses proteins; cysteine proteinase c; Superfamily: papain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession: T10518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 20-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Species: Ananas comosus (pineapple)
                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Experimental source: cv. N67-10; nearly mature fruit
                                                                                                                                                                                                              123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               132 LPESIDWREKGAVTAVKNOG 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96 VPQSIDWRDYGAVNEVKNQ 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 VPQSIDWRDSGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                              1 VPQSIDWRDSGAVTSVKNQ 19
                                                                                                                                                                                                              VPQSIDWRDYGAVNEVKNQ 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                     84.21;
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                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                               Score 84; DB 2; Les
Pred. No. 4.8e-06;
Pred. No. 4.8e-3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 4.4e-06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 351
                                                                                                                                                                                                                                                                                                                                                                                        Indels
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C.Superfamily: papain
C.Superfamily: papain
F:1-17/Domain: signal sequence #status predicted <510>
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                                                                                                                                                        A;Pathway: protein degradation
A;Note: important role in the lysosomal degradation of proteins
                                                                                                                                                                                                                                                                                                       C/Function
                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: protein
A; Residues: 18-31 < OK2>
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A.Experimental source: epididymis
A.Experimental source: epididymis
A.Experimental source: epididymis
A.Experimental source: and CCC for residue 203 as Thr and CCC for residue
A.Experimental source: and CCC for residue 203 as Thr and CCC for residue
A.Experimental source: and CCC for residue 203 as Thr and
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Fil-26/Comain: agundl sequence stratus predicted <810>
Fil-26/Comain: agundl sequence stratus predicted <800<
Fil-27-134/Comain: propeptide stratus predicted <800<
Fil-27-134/Comain: propeptide stratus predicted <800</li>
Fil-27-134/Comain: propeptide stratus predicted <800</li>
Fil-27-134/Comain: propentide stratus experimental files
Fil-27-134/Comain: propentide stratus experimental files
Fil-27-134/Comain: propentide stratus experimental files
                                                                                                                                                                                                                                                  A Description: catalyzes hydrolysis of peptide bonds in proceins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Accession: S5991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A/Cross-references: DDBJ:D37917; NID:g710655; PIDN:BAA07140.1; PID:g1468964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A,Accession: A58195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C, Superfamily: papain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;159,293,313/Active site: Cys, His, Asn #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        :Residues:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              136 PQSIDWRAKGAVTPVKNQG 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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6. Similarity

Conservative

84.2%; Pred. No. 4.8e-06; 0 Mismatches

Length 352 3; indeis

c Gaps

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TO9760

Lysosome; procein

chain

precurso

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Systems, DCC; Magneth, M; LVMD, K.R.
BLOCHDE, UZ 266, 7981, 3990
A-TICLE: TIM mittop ocid sequence of Chypopapain from Comitoe papaya.
A-TICLE: TIM mittop ocid sequence of Chypopapain from Comitoe papaya.
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C.Species Carte papaya (papaya)
C.Date: 16-UDI 1999 Beegruence revision 16-UDI-1999 #rext_change 18-Aug-2000
C.Accession: (1997-65), S04225, S06285
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A; Accession: T09760
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F.18-113/Domain: propeptide Stratus predicted <PRO.
F.18-113/Domain: propeptide Stratus predicted <PRO.
F.18-178,169-272,720-232)/Dismitée bonds: stratus predicted
F.18-178,104/Accider: site: Oys. His. Ams instanus predicted
F.18-272,222/Binding site: carbohydrate (Asm) (covalent) stratus predicted

Gaps

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79.0%; Score 83; DB 2; Length 334; 70.0%; Pred. No. 6.7e-06; 2ve 4; Mismatches 2; Indels

Best Local Similarity 70.09

Ouery Match

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Rightment V. Astria, v. Astria, v. Assers, v. As on Research, J.; Schnek, A.G.; Risinschmidt, T.; Loose, Y. Marisen Chem. Mapper-appler 774; 503-566, 1993.
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C.Speckes Carlos puescens (unountain papaya)
C.Speckes (0-14)94 #sequence_revision 01.Nov-1
Gaps
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3; Indels
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A.Map position: 5
A.Introns: 34/3; 102/3; 140/2; 180/2; 336/2; 357/2; 417/1
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Pred. No. 1e-05;
Mismatches
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Anderciption: A movel sulthydryl endOpeptidase is induced by environmental stress and A/Reference number: 227502
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cysteine proteinae (BC 3.4.12.-) - common ice plant Chicariata namae: oyecine deooppeptdaes allthydyy-endopptidae C;Species, Mesembryanthemm crysteilinum (common ice plant) Cpiccession: 171373 Setquence\_revision 23.401-1599 Hext\_thange 04-Peb-2000

79.0%; Score 83; DB 2; Length 367; 80.0%; Pred. No. 7.4e-06;

Segt Local Similarity Superfamily: papain

C.Keywords: cysteine proteinase, hydrolase F;150,287,308/Active site: Cys, His, Asn #status predicted

Gene: Sep1

Genetics

A. Accession: T1358
A. Accession: T1358
A. Molecule: pre-liminary:
A. Molecule: pre-liminary:
A. Molecule: pre-liminary:
A. Molecule: pre-liminary:
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A. A. Cross Enferonce: Leave Mol. Strossed; closs pSepp.
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Gaps

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Length 357;

Score 83, DB 2, Length 357 Pred. No. 7,2e-06; 1, Mismatches 1, Indels

79.0%;

Query Match

16; Conservative Best Local Similarity

Matches

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RESULT 13

C.Keywords: cysteine proteinase; hydrolase Fil-Z6/Domain: signal sequence Heactus predicted <81G> Fil-26/Domain: signal sequence Heactus predicted <84MI> Fil47,281,302/Active site: Cys, His, Asn Heacus predicted

Superfamily: papain

A/Status: preliminary; translated from GB/EMBL/DDBJ A/Molacule type: mRNA A/Residues: 1-357, <RGB> A;Cross-references: EMBL: AJ009830; PIDN: CAA08861.1

A:Accession: T07851

ananain (EC 3.4.22.31) precursor AN11 - pineapple

RESULT 12

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	100 0 100 0 10	. Is the number of results predicted by chance to have a sater than or equal to the score is the result being printed entryed by analysis of the coral score distribution.  SUMMARIES  OURTY MARCO Length DB ID  Description	Mariam De eq langh: 200000000 Mariam De eq ingdh: 200000000 Por-processing: Mariam March 0% Forther Statum March 0% Listing first 45 summaries Database: 1.07.71. 2.07.71. 3.07.71. 4.07.71. 4.07.71.	Searched: 283224 seqs, 96134422 residues Total number of hits satisfying chosen parameters: 283224	NOTION: NOVEMBER 21, 2002, 161714 () Dearch Fine 13 Seconds  NOTION () Dearch Fine 13 Seconds  147.899 Million call updates/sec  118.09-674-718-1  Sequence: 108.09-674-718-1	ein - pro
AuStatus preliminary translated from GB/BBML/DBMJ Albacated PTPS cmM3 Alcost reterances (BBML/DBMJ) Alcost reterance (BBML/DBMJ) C. Specifically, papeals reterance attack predicted cfGO F. SCHOOL (BBML/DBMJ) F. SCH	RESULT 2  RESULT 2  PIGGA: Be seen becombain (DC 3 4.22.23) precursor - pineapple  FIGAL terms result produces  C.Species: Annas comonus (pineapple)  C.Dee; 15.201.2399 Seconter_revision 15.01.1399 Hear_change 20-Jun-2000  C.Dee; 15.01.2399 Seconter_revision 15.01.1399 Hear_change 20-Jun-2000  R.Mate, Be; Odensolo, Y; Ota, S.  Albertopion; Coning and sequencing of cysteine proteinses in Annas comonus.  A.Accession; Tiblés allos Sequencing of cysteine proteinses in Annas comonus.  A.Accession; Tiblés allos Sequencing of cysteine proteinses in Annas comonus.	C:Reywords: cyneann proteinae; hydrolase Coary Macchae; billot #; Boog 168; BB 2; Length 212; Bent Local Shilarity, 100 6%; Bred: Bo 7; 18-10; Macchae; 20; Comervative 0; Mismatches 0; Indels 0; Gaps 0; Cy 1 vPgstrokorkarisywowa 20 1                  Db 2 vPstrokorkarisywowa 21	RESULT 1  RESULT 1  RESULT 1  RESULT 1  RESULT 2  RESULT 2  RESULT 3  RESULT	ALIGNENTS	3 71 68 7 61 1 National Organia Nutra 24 1 1 National Organia Nutra 24 1 National Organia Nutra 24 1 National N	73 67.6 317 2 84451 72 66.7 331 2 UCS441 72 66.7 331 2 UCS441 72 66.7 331 2 UCS441 72 66.7 351 2 UCS441 72 66.7 361 2 UCS451 72 66.7 361 2 UCS451

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§ Kapper, A.D.; Bornett, S.P.; Horowski, M.; Holderidge, M.B.; Jeanard, M.J.C.; Rogers, B. Borders, C. B.; A. 101, 727, 738, 7394.
Reserve the state of material and adversariation of multiple forms of the pineapple-steme-denive. A Accessing Section 1982 50, 948-94, 9410-14319-94, PMIL-8539-8
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                C;Species: Anamas comosus (pineapple)
C;Date: 16-Jul-1299 #sequence_revision 16-Jul-1399 #text_change 20-Jun-2000
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C.Date: 16-Peb-1995 #sequence_revision 01-Nov-1996 #text_change 07-May-1999
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F)-14/10main: asgmi sequence status predicted GSIOs

F)-15/11/10main: asgmi sequence status predicted GROs

F)-12/10/10main: asmooning terminal propegated seature predicted GARS-

F)-10/10/10main: carbooning terminal propegation status predicted GCTP-

F)-10/10/10/Main: carbooning cerminal seature predicted GCTP-

F)-10/10/10/Main: carbooning attention and seature predicted
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A,Experimental source: cv. N67-10; nearly mature fruit
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C.Keywords: cysteine proteinase, hydrolase
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A,Residues: 1-20 <MAP>
C,Superfamily: papain
C,Keywords: cysteine proteinase; hydrolase
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A, Residues: 1-20 <NAP>
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A, Residues: 1-351 <MUT>
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Ributa, E.; Odestock, Y.; Otta, S.
Albertigted to the Bolks Data Library, October 1994
Albertiption: Closing and sequenting of cysteline proteinases in Annass compans.
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Alberti
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C.Speckes: Amanas comouns (pineapple)
C.Speckes: 16-UJL-1999 Hecquenc (pineapple)
C.Dates: 16-UJL-1999 Hecquence, revision 16-UJL-1999 #text_change 20-JUN-2000
                                                                                                                                                                                        0; Gaps
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19-313/Podutt: fruit bromelain #stetus predicted «NMT»
P.334-324/Omain: actoryvi-terminal propeptide #stetus predicted «CTP»
P.304-324/Omain: actoryvi-terminal propeptide #stetus predicted (CTP»
ch 100.0%; Score 108; DB 2; Length 356; 15:milarity 100.0%; Pred. No. 1.38-09. 113-09. 1100.0%; Pred. No. 1.38-09.
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C.Neywords. Cysteine proteinase; hydrolase
F;148,360,30,74ctive sice: Cys, H3s, Asn #status predicted
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A;Experimental source: cv. N67-10; nearly mature fruit
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A;Molecule type: mRNA
A;Residues: 1-324 <MUT>
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Duery Match Best Local S

9

Matches

Genetics

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C.Superfamily papain
C.Reywords: cysteine proteinass; hydrolass
F:1-26/Domain: Signal sequence Hetatus predicted cSIG>
F:27-357/Product: anamain MAN1 Hatatus predicted cMAT>
F:147,281,302/Active Site: Cys. His, Asn Hetatus predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Robertson, C.B.; Goodenough, P.W. submitted to the RMSL Data Library, July 1998 submitted to the RMSL Data Library, July 1998 A.P.Becription: Coning and expression of anamain gene from pineapple. A.Reference number: 216167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ananain (E. 3., 4.2.31) procursor AN1: primappie
W.Alternate manes; cynteine procedanae precursor AN1:
C.Species: Ananae commons (pinéappie)
C.Species: 14-My.1993 Respense, excision 14-May-1999 West_change 20-Jun-2000
C.Mccession: T0935 Condenough, P. M.
M.Moberson, C. B. W. Condenough, P. M.
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C.Superfamily: papain
C.Superfamily: proteinase: hydrolase
C.Superfamily: proteinase: hydrolase
C.Superfamily: hydrolase: hydrolase
C.Superfamily: hydrolase: hydrolase
F;147,279,100/Active site: Ope.
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CJGenetics:
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A;Molecule type: mRNA
A;Residues: 1-345 <ROB>
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submitted to the EMBL Data Library, November 1997
h.Description: Cloning and expression of ananain gene from pineapple
h.Reference number: 216166
h.Rocession: 707839
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A;Status; preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-357 <ROB:
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C/Species: Ananas comosus (pineapple)
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                      123 VPQSIDWRNYGAVTSVKN 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VPOSIDWRDYGAVTSVKNO 19
                                                                             1 VPQSIDWRDYGAVTSVKN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
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                                                                                                                                     conservative
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                                                                                                                                                           94.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         86.1%; Score 93;
94.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                           Pred. No. 5e-07; Length 357;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 1.4e-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 93;
                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OB 2; Length 345;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
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C.Reywords: Cysteline proteinass; hydrolase
Fl-24/Domain: asignal sequence stratus predicted cRIG>
Fl-25-121/Domain: asimo-terminal prespected stratus predicted cPRO>
Fl-25-239/Product: fruit bromelain Basatus predicted cMAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-340 <MUT>
A; Cross-references: EMBL: D38533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A.Description: Cloning and sequencing of cysteine proteinases in Ananas comosus.
A.Reterence number: 217065
A.Accession: 110516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Muta, E.; Okamoto, Y.; Ota,
submitted to the EMBL Data Lil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fruit bromelain (EC 3.4.22.33) FB22 precursor - pineapple (fragment)
C:Species, Ananas comosus (pineapple)
C:Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 20-Jun-2000
                                                                                                                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-352 MUTS
A; Cross-references: EMBL:D14058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A.Description, Cloning and sequencing of fruit bromelain
A.Paference number: 217060
A.Accession: T19501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fruit hromelain (EC 3.4.22.33) BB13 precursor - pineapple
C:Spocies: Anahase comosus (pineapple)
C:Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 20-Jun-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Function:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary; translated from GB/EMBL/DDBJ
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cysteine proteinase (EC 3.4.22.-) - Douglas tir
                           95850
                                                                                                                                                                                                                                                                                                                                                             F;148,281,302/Active site: Cys, His, Ash #status predicted
                                                                                                                                                                                                                                                                                                                                                                              F.1-24/Domain: signal sequence stratus predicted «510»
F.25-122/Domain: asino-terminal propelled status predicted «ART»
F.123-130/Froduct: Eruit bromelain Status predicted «ART»
F.131-135/Domain: carboxy-resminal propeptide status predicted «CTP»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Keywords: cysteine proteinase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C.Superfamily: papain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Description: hydrolyzes proteins; cysteine proteinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Experimental source: cv. N67-10; rrult; nearly mature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: translated from GB/EMBL/DDBJ
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submitted to the EMBL Data Library, January 1993
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                                                                                                                                                                                                                                                                          83.3%; Score 90; DB 2; Length 352
85.0%; Pred. No. 1e-06;
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85.0%; Pred. No.
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FF.13-14/Domain, propeptide seasus predicted «Sto-
FF.135-146/Product: operation proteinase Sto-
FF.155-146/Product: operation proteinase Sto-
FF.155-142/Product: operation proteinase Producted «PAT»
FF.155-142/Productive stice operation predicted «PAT»
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                                                                                                                                                                                                                                                                                                                                        73.1%; Score 79; DB 2; Length 364;
73.7%; Pred. No. 6.3e-05;
Live 2; Mismatches 3; Indels
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Pred, No. 9.3e-05;
3; Indels
Avintrons: 148/1, 274/3
Cykoptefamily: papain
C;Koywords: cysteine proteinase, hydrolase
P.150,287,308/Active site: Cys, His, Asn #status predicted
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Job time : 14 secs
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                                         C Species: Standard naw momental (1004) 100-01-1916 Steak_change 04-Peb-2000
C Appellant Standard naw momental (1004) 100-01-1916 Steak_change 04-Peb-2000
C Appellant Standard Name C Standar
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C;Date: 07-Jul-1995 #sequence_revision 07-Jul-1995 #text_change 20-Jun-2000
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C. Appending Properties properations by Montal apidocale C. S. Speciale properations by Montal apidocale SP144, 588, 196 Apidocale safety Cys. His. Ann Heneus predicted
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C.Keywords: cysteine proteinase; hydrolase
F.156,292,312/Active site: Cys. Ris. Asm #status predicted
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A,Map position: 3
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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             length: 2000000000
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11 /ogn2 6/pcodata/1/aa/5A_COMB.pop:

22 /ogn2 6/pcodata/1/aa/5B_COMB.pop:

31 /ogn2 6/pcodata/1/aa/6B_COMB.pop:

41 /ogn2 6/pcodata/1/aa/6B_COMB.pop:

51 /ogn2 6/pcodata/1/aa/6B_COMB.pop:

52 /ogn2 6/pcodata/1/aa/backf1les1.pop:

63 /ogn2 6/pcodata/1/aa/backf1les1.pop:
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Copyright (c) 1993 - 2002 Compugen Ltd.
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                                US-08-578-701A-1
US-08-360-693-1
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US-09-325-932A-167
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sequence 17, Applisequence 18, Applisequence 18,
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; Sequence 1, Application US/08578701A
; Patent No. 5824305
                                                                                                                                                                                                                                                                                                                                         US-08-578-701A-1
                                                                                                                                                                                               Query Match 100.0%; Score 108, DB 2;
Best Local Similarity 100.0%; Pred. No. 9-5e-11;
Matches 20; Conservative 0; Mismatches 0;
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CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION: 18

NAME: Baldwan, Geraldine F. REDISTRATION MUNBER: 31.22

REDISTRATION MUNBER: 31.212

TELECHMONICATION INFORMATION: TELECHMONICATION INFORMATION:
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TYPE: amino a
STRANDEDNESS:
TOPOLOGY: unk
MOLECULE TYPE:
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: MEDICAL USE OF STEM BROWELAIN PROTEASE
NUMBER OF SEQUENCES: 1
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MEDIUM TYPE: TJOPPY disk
COMPUTER: TIPM PC compatible
OPERATING SYSTEM: PC-DSC/MS-DOS
SOPTHARE: PALENTIN BELEASE #1.0, Version
APPLICATION NUMBER: US/08/$78,701A
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STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
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                                                                                                                                   1 VPQSIDWRDYGAVTSVKNQN 20
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81.54; Score 88; DB 4; I
Best Local Similarity 88.94; Fred. No. 1.1e-08;
Best Local Similarity 88.94; Stred. No. 1.1e-08;
Best Local Similarity 80.94; Stred. No. 1.1e-08;
CHRRANT FLIND MATR: 199-06-25

PRICRA PAPELLOY ON NAMERS: CT/CR89/00552

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CURRENT FILING DATE: 1999-06-04
UNDREN OF SEQ ID NUS: 266
SOFTWARE: FRRESEQ FOR WINDOWS VERSION 3.0
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Patent No. 6451604
GENERAL INFORMATION:
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Patent No. 6451604
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Best Local Similarity 68.49
Marches 13, Conservative
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US-09-382-689A-1
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US-09-325-932A-167
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100.0%; Pred, No. 9.5e-11;
tive 0; Mismatches 0; Indels (
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WEDIUM YNEE: Rioppy dark
COMPUTER: READERS FORM;
COMPUTER: REA
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3000 K Street, N.W., Suite 500
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FILING DATE: 30-JUN-1993
PRIOR APPLICATION DATA:
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PELIND DATE: A5-UNI-1993
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PELINDATION NUMBER: CB 930184-4
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PELINDATION 
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FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
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Patent No. 6335427
GENERAL INFORMATION:
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APPLICANT: Engwerda. Christian
APPLICANT: Peek, Keith
TITLE OF INVENTION: Component of
FILE REFERENCE: 0623.0760031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPHAX: (202)672-539
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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US-09-382-689A-1
               RESULT 2
US-08-360-693-1
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APELCOMY: MARCET, Shunji APELCOMY: ARCET, Shunji APELCOMY: ARCET, Shunji APELCOMY APPLICATION APPLICAT
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; ORGANISM: Eucalyptus grandis
US-09-325-932A-165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-325-932A-165
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; Sequence 80, Application US/09120365 ; Patent No. 6103514
                                                                                                                              RESULT 8
US-09-120-365-80
                                                                                                                                                                                                                                                                                                                                                                    ğ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-120-365-62
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Patent No. 6103514
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 102/09/325,932A, CURRENT APPLICATION NAMERS: US/09/325,932A, CURRENT FILLING DATE: 1999-06-04 NAMERS OF SEQ ID NOS: 208 SEQ ID NOS: 208 SEQ ID NOS: 208 SEQ ID NO 108 SE
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Patent No. 6451604
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APPLICANT, Leabna, Annete to Compositions affecting programmed cell TITHS OF INVESTIGATION: Compositions affecting programmed cell TITHS OF INVESTIGATION. Compositions and their use in the modification of forestry plant development of INVESTIGATION CONTROL OF T
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ORGANISM: Sarcophaga cathepsin L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     144 LPEDFDWRDHGAVTGVKNQ 162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      57 LPKDFDWRDHGAVTAVKNO 75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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68.4%; Pred. No. 5.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73.1%; Score 79; DB 3; Length 339; 68.4%; Pred. No. 1.5e-05; ative 5; Mismarches 1; Indels
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Pred. No. 1.2e-05;
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US-09-120-365-80
                                  ; ORGANISM: Sarcophaga cathepsin L
US-09-515-039-80
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REQUISION 80. Application US/09515039

PREDER No. 62.1559

GENERAL INCOMPATION SHUT!

TITLE OF INVESTIGN INV PROTENSE

TITLE SEESENGE 1230-14449

GENERAL REPORT OF THE SEESENGE 1230-1449

GENERAL RE
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CURRENT PELLICATION NUMBER: US/09/515,039

CURRENT PELLICA DATE: 2000-03-06

SALLER APELLICATION NUMBER: 19-333 474

BALLER FILING DATE: 1997-11-18

NUMBER OF SEQ 1D MOS: 101

SOFTMARE: Referent Number: 2.0

SOLUTION 63.

SOLUTION 63.

SOLUTION 63.
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US-09-515-039-62
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Sequence 62, Application US/09515039
Patent No. 6214599
GENERAL INFORMATION:
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CURRENT APPLICATION HYMNER: US/09/120,365
CURRENT FILING DATE: 1996-07-22
ENNLIER APPLICATION HYMNER: JP 9-333 474
ENNLIER FILING DATE: 1997-11-18
MOMBER: 09 580 1D NOS: 101
NOMER'S PAGE 17 101
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TITLE OF INVENTION: NEW PROTEASE
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Pred. No. 1.5e-05;
5; Mismatches 1
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Pred. No. 1.5e-05;
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Best Local Similarity 68.4%, Pred. No. 9.9e-05.
Matches 13, Conservative 3, Mismatches 3, Indels
PAULICANT, NOCTI, SHOLLS, TITLE OF JUNEARITON HER RECEIVED OF CHEERER RECEIVED AND HER RECEIVED OF CHEERER RELIES OF CHEERER FELLING DATE, 1998-07, 22 CHEERER FELLING DATE, 1998-07, 22 CHEERER FELLING DATE, 1998-07, 22 CHEERER FELLING DATE, 1999-07, 22 CHEERER FELLING DATE, 1997-11, 18 CHEERER PROPERTIES OF CHEERER PROPERTY PROPERTIES OF CHEERER PROP
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CURRENT APPLICATION NUMBER: US/95/15,019
CURRENT FILING DATE: 2000-05-06
EMBLIER APPLICATION NUMBER: UP 9-333-474
EMBLIER FILING DATE: 1997-11-18
NUMBER FILING DATE: 1997-11-18
SUPPLIER FILING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 64, Application US/09120365; Patent No. 6103514; GENERAL INFORMATION:
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TITLE OF INVENTION: NEW PROTEASE
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Matches 13; Conservative
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US-09-120-365-64
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SEQ ID NO 64
LENGTH: 334
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US-09-120-365-69
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US-09-120-365-64
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                                                Ouery Match 73.1%; Score 79; DB 4; Length 339; Best Local Smilarity 60.4%; Pred. No. 1.56-65. Matches 13; Conservative 5; Mismatches 1; Indels
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COMPUTER. IBM Compatible
OFFEATING SYSTEM. DOS
COMPUTER. PRESENCE for Windows Version 2.0
CURRENT APPLICATION UNDAY.
APPLICATION UNDAY.
FILING DATE: HEFFAILE.
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Sequence 4, Application US/OBESISSE
APPLICATION Enthusing Objective Application of Application Objective Application of Application Objective Application Objective Application Objective Application Objective Application Objective Application Objective Objectiv
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, Patent No. 6103514
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REFERENCE/DOCKET NUMBER: PF-
TELECOMMULICATION INFORMATION:
TELEPHONE: 415-855-0555
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SEQUENCE CHARACTERISTICS:
LENGTH: 334 amino acids
TYPE: amino acid
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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MEDIUM TYPE: Diskette
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CLONE: GenBank
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US-09-120-365-69
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Quary Match
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Sectional Similarity (6.5): Score 74; D8 3; Length 34;
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Sectional Similarity (7.7)
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Result
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Maximum Match 100*
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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(without alignments)
51.170 Million cell updates/sec
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11 / 1/9072 6/pcdata/1/isa/5A_COMB_pep:

22 / 1/9072 6/pcdata/1/isa/5B_COMB_pep:

32 / 1/9072 6/pcdata/1/isa/6E_COMB_pep:

33 / 1/9072 6/pcdata/1/isa/6E_COMB_pep:

44 / 1/9072 6/pcdata/1/isa/6E_COMB_pep:

55 / 1/9072 6/pcdata/1/isa/backfiles1.pep:

66 / 1/9072 6/pcdata/1/isa/backfiles1.pep:
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Copyright (c) 1993 - 2002 Compugen Ltd.
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; Sequence 1, Application US/08578701A
; Patent No. 5824305
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  US-08-578-701A-1
                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 213 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: unknown
MOLECULE TYPE: protein
                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CHRENT APPLICATION DATA.
CHRENT APPLICATION DATA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10036
COMPUTER READABLE FORM
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Mynott, Tracy L.
TITLE OF STEM BROMELAIN PROTEASE
NIMBER OF SEVENTION: MEDICAL USE OF STEM BROMELAIN PROTEASE
NIMBER OF SEVENTENCES:
ADDRESSORDENCE ADDRESS:
ADDRESSORDENCE ADDRESS:
ADDRESSORDENCE ADDRESS:
ADDRESSORDENCE ADDRESS:
ADDRESSORDENCE ADDRESS:
ADDRESSORDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Baldwin, Geraldine F.
REGISTRATION NUMBER: 31,232
REFERENCE/DOCKET NUMBER: 71:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
FILING DATE: 22-DEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                    1 VPQSIDWRDSGAVTSVKNQ 19
                                                    VPQSIDWRDYGAVTSVKNQ 20
                                                                                                                                                        Conservative
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                                                                                                                                                                              94.7%
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US-08-751-102-2
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US-08-
                                                                                                                                                        0;
                                                                                                                                                                                   Score 93;
Pred. No. 1
                                                                                                                                                        Mismatches
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                                                                                                                                                                                   .1e-07
                                                                                                                                                        1; Indels
                                                                                                                                                                                                        Length 213;
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sequence 11. Appli
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sequence 13. Appli
sequence 13. Appli
sequence 14. Appli
sequence 15. Appli
sequence 15. Appli
sequence 15. Appli
sequence 15. Appli
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Gaps 0 Page 2

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  79.0%; Score 83, DB 3; Length 334; 70.0%; Pred. No. 7.8e-06; 2: Indela
                                                                                                                                                         MEDIWAY WYSE. Diskette
COMPUTER. LBM Compatible
CHEMPING SYSTEM. DOS SERVING STEM.
CREATER RESERVED TO WINDOWS Version 2.0
CREATER APPLICATION WORN.
APPLICATION WORSE. US/08/883,526
FILING DATE: Herealth.
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRICIACHY NACCI.; SUNIA
PRICIACHY NACCI.; SUNIA
CHERAFRESHICH SHARP POUTBAS
CHERAF PELINO DATE: 1995-07-22
CHERAF PELINO DATE: 1995-07-23
PARIER APPLICATION NAMES: UP -233 474
NAMES NE LINO DATE: 1997-11-18
SUNIAGE PECENTIN NACCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                    PF-0331 US
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Patent No. 6103514
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-0555
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    114 VPKSVDWREKGYVTAVKNOG 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : 334 amino acids
amino acid
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single
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CLONE: GenBank
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                                                          Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
                                                                                                  TIGA
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                                                                                                                 94304
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US-09-120-365-69
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                                                                                                  COLDUTRY
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                                                          CITY: P
STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                  APPLICANT: MYNOTT, Tracey L. Tracey E. TILLE OF INVENTION: 18S OF ENZYMES, ESPECIALLY BROMELAIN, IN TITLE OF INVENTION: THE TREATMENT OF DIABRHOEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 88.6%, Score 93; DB 2; Length 213; Best Local Smilarity 94.7%; Pred. No. 1.1e-07; Indels Matches 18; Conservative 0; Mismatches 17 Indels
                                                                                                                                                                                                                                                                                                              COMPUTER EAGLBLE FORM:
MEDITUR TYPE: Floppy disk
COMPUTER: IRM FCompatible
OPERATING SYSTEM: FC-DOS/NS-DOS
SOFTWARE: Featuring Figure 6.005/NS-DOS
COMPARING Peaturing Figure 6.005/NS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30357/115/KIST
                                                                                                                                                                                            S: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bandman, Olga
APPLICANT: Godgler, Karl J.
APPLICANT: COCIey, Neil C.
APPLICANT: Stab, Purvi
TITLE OF INVENTION: NEW HUMAN CATHEREIN
NUMBER OF SEQUENCES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/GB93/01374
FILING DATE: 30-JUN-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/360,693
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
                                       Sequence 1, Application US/08360693
Patent No. 5928640
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/08883526
Patent No. 6033893
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMOUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 VPQSIDWRDYGAVTSVKNQ 20
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 904136
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & L
                                                                                                                                                             NIMBER OF SECRENCES.
                                                                                                                                                                                                                                       Washington
                                                                                                                                                                                                                                                                                                  20007-5109
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                   US-08-360-693-1
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                                                                                                                                                                                                                                                           STATE:
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US-09-325-932A-176
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US-09-515-039-69
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                                                                                                                                                                                                                                                                                                                 US-09-325-932A-167
                                                                                                                                                                                                                                                                                                                                                 KESOLT /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-U9-325-932A-176
                                                                                 OMERGAL PROPARTIES
APPLICANT: Labour, Admits
APPLICANT
A
                                                                                                                                                                                                                                             Sequence 167, Application US/09325932A
Patent No. 6451604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 176, Application US/09328932A
Patent No. 6451604
Patent No. 6451604
APPLICANT: Filin, Barry
APPLICANT: Lasham, Annette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PATENT NO. 6214599
GENERAL INFORMATION:
APPLICANT: Natori, Shunji
TITLE OF INVENTION: NEW PROTEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/325,932A
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEC ID NOS: 206
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CURBENT PALLACTION NUMBER US/09/515,039

CURBENT PILLNO DATE. 2000-03-06

BARLIER APPLICATION NUMBER, UP 9-333 474

BARLIER FILMS DATE. 1997-11-8

NUMBER OF SEQ 10 NOS: 1017-18

SOFTWARE OF SEC 10 NOS: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 69, Application US/09515039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lasham, Annette
ITILS OF INVENTION: Compositions affecting programmed cell
ITILS OF INVENTION: death and their use in the modification of forestry plant devel
ITILS SERENCE: 1022
CURRENT APPLICATION NUMBER: US/09/325,932A CURRENT FILING DATE: 1999-06-04
                                                                    KILE KEFEKENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Eucalyptus grandis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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:|: |||| ||||:|||||
67 LPKDFDWRDHGAVTAVKNQG 86
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70.0%; Pred. No. 1.1e-05;
mtive 3; Mimmatches 3; Indels
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US-09-325-932A-167
                                                                                                                                                                                                                                                   ; ORGANISM: Murine cathepsin L
US-09-120-365-64
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US-09-325-932A-165
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SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                           GEREBAL INCOMMOTION:
APPLICANT NACCH, Shunyi
TTILE OR INVENTION; NEW PROTEASE
FILE REFERENCE: 12280-144749
CURRENT APPLICATION NUMBER: US/09/120,365
CURRENT FILMS DATE: 1398-01-22
SABLIER APPLICATION NUMBER: UP-9-33 474
BALIER FILMS APPLICATION NUMBER: UP-9-33 474
BALIER FILMS APPLICATION NUMBER: UP-9-34 474
BALIER FILMS APPLICATION NUMBER: UP-9-34 474
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Patent No. 6103514
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SEQ ID NO 167
LENGTH: 148
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Best Local Similarity
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Patent No. 6451604
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      114 IPKSVDWREXGCVTPVKNQG 133
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65.0%; Pred. No. 1.1e-05;
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Pred. No. 4.3e-06;
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TITLE OF INVARIATION USE PROFILES

TITLE OF INVARIATION USE PROFILES

GREEN FILES APPLICATION PROSES US/05/515.03

CREENER FILES OF SEC. 2006-63-06

CREENER FILES OF SEC.
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Patent No. 6214599
GENERAL INFORMATION:
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US-09-120-365-80
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ORGANISM: Sarcophaga cathepsin L
US-09-515-039-62
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US-09-515-039-80
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SEQ ID NO 80
LENGTH: 339
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 80
LENGTH: 339
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Best Local Similarity
Matches 14, Conserv
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US-09-515-039-62
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Best Local Similarity 65.0%; Pred, No. 1.1e-05;
A. Mismacches 3; Indels
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TITLE OF INVENTION, NEW PROCESSE
CHERENT APPLICATION MOREIN. US/0.9/120,365
CHERENT APPLICATION MOREIN. US/0.9/120,365
EMBLIER FAILTON OF MOREIN. US/0.9/11-18
EMBLIER FAILTON DATE: 1997:11-18
SOFFWARE OF REAL TO NOS: 1.0
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CURRENT FILMS OFFER: 3000-03-6
SARLIER APPLICATION NOMBR: UF 333 474
RALLIER FILMS OFFE: 1997-11-18
WOMBER OF SED 1D NOS: 101
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TITLE OF INNERTION NEW BEOTENSE
CHERENY ADDITION DATE: 1929-17.25
CHERENY THEORY DATE: 1929-17.25
EARLIER FILEND DATE: 1929-17.19
EARLIER FILEND DATE: 1929-11.19
NUMBER OF SEQ IN NOS: 101
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Sequence 64, Application U8/09515039
Patent No. 624599
GENERAL INCRMATION: Bunji
TITLE OF INVENTION: NEW PROTENSE
FILE REPERENCE: 32390-144749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 62, Application US/09120365
Patent No. 6103514
GENERAL INFORMATION:
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US-09-515-039-64
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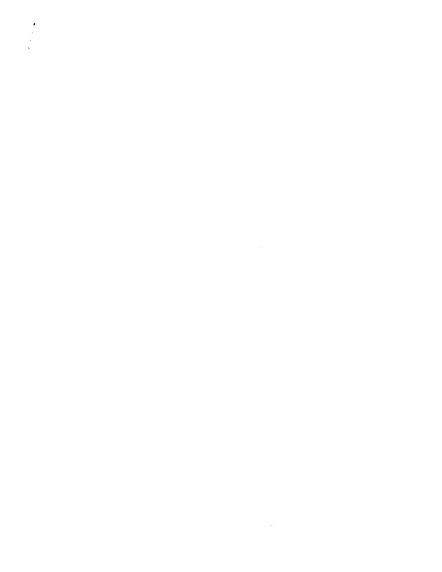
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Search completed: November 21, 2002, 16:11:01 Tob time: 11.5 secs
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Best Local Similarity 70.0%, Pred, No. 2e-05;
Matches 14; Conservative 3; Mismatches 3; Indels
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Maximum Match 100%
Listing first 45 summaries
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Match
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108
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Copyright (c) 1993 - 2002 Compugen Ltd.
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20 US-09-674-738-1

17 US-09-382-688-3

17 US-09-382-688-3

21 US-09-750-210A-3

6 US-08-244-796-15

21 US-09-791-537-84879
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Sequence 1, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 15, Appl
Sequence 84879, A
                                                                                                                                                                                                     Description
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GENERAL INFORMITION:
APPLICANT URSAPHAN Arrainite1 (mbH
TITLE OF INVENTION: Use of Encemains Proteases for Inhibiting Slood
TITLE OF INVENTION: Compilation
FILE REFERENCE, 10054
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US-09-674-738-1
                                                        1 VPOSIDWRDYGAVTSVKNON 20
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-09-513-996A-9	09-708-427-2093	-10-219-220-16	90-004A-1	-09-327-373-167	-10-219-220	-990-004A-1	-09-327-373-17	-620-393B-334	-09-620-393B-33	-09-791-537-3225	:-09-791-537-119E	-10-029-301-	-09-750-210A	-09-743-987-2	-382-688-2	-09-791-537-6933	-09-791-537-1107	9-791-537-8	-09-791-537-8486	-09-791-537-8240	-09-750-210A-	-09-674-73	-09-382-688-	-09-382-	-09-743-987-1	-09-743-987-	-09-743-98	-09-743-987-	-09-743-987-2	-09-750-2102	-09-382-688	-09-382-685A-4	-09-791-537-	-09-743-987-	-09-791-537-6	-09-743-987-2	-09-743-987-16	9-791-5
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Qy 1 VPQSIDMRDYGAVTSVKNQN 20	Overy March 100.04; Score 108; DB 20; Length 20; Sect Local Similarity 100.04; Pred Mo 6 Sec.10; Indels 0; Gaps Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps	ORCANISM: pine-apple (Bromeliacea) US-09-674-738-1	SEQ ID NO 1 LENGTH 20	; NUMBER OF SEQ ID NOS: 2 ; SOFTWARE: PatentIn Ver. 2.1	PRIOR APPLICATION NUMBER: PCT/EP98/04406 PRIOR FILING DATE: 1998-07-15	CURRENT FAILING DATE: 2000-10-31

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15-09-382-685A-2 Sequence 2, Application US/09382685A GENERAL INFORMATION:

PRIOR APPLICATION NUMBER: GB 9706119.6 PRIOR APPLICATION NUMBER: GB 9706119.6 PRIOR APPLICATION NUMBER: GB 9704252.7 PRIOR FILING DATE: 1997-02-28 PRIOR FILING DATE: 1997-02-28

PRIOR INCLUSION NUMBER GB 9703827.7 PRIOR FILMED DATE: 1997-02-25 NUMBER OF SEQ ID NOS: 4
SOFTWARE PARENTIN VET. 2.0 TYPE: PRT ORGANISM: ORGANISM: Ananas comosus

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APPLICANT: ANDERN R. MILMER, MICHAEL PANACCIO, TERRNCE W.
MILTE OF INVESTIGEN. LIVER FLUKE VACCINE AND POLYBEFTIDES USEFUL FOR SAME
NUMBER OF SEQUENCES: 24
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ADDRESSEE: SCULLY SCOTT MURPHY & PRESSER
STREET: 400 GARDEN CITY PLAZA, GARDEN CITY
                                                                                                                                                                                                           APPLICANT: Winott, Tracy Lehanne
APPLICANT: Bouwerd Christian
APPLICANT: Edward Christian
APPLICANT: Peek, Keith
TILL OF INVENTION: Component of Exomelain
FILE REPRENCE: 0623.0740001
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11530-0299
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US-09-750-210A-3
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Sequence 15, Application US/08244796
GENERAL INPORMATION:
                                                                                       Sequence 3, Application US/09750210A
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2 VPOSIDWRDYGAVTSVKNON 21
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APPLICATION NUMBER: PCT/A
FILING DATE: 04-FEB-1994
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: NEW YORK
STATE: NEW YORK
                                                  RESULT 4
US-09-750-210A-3
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Query Match 100.0%; Score 108; DB 17; Length 21; Best Local Similarity 100.0%; Pred. Mo. 7.7.3-0.7 Marches 20; Conservative 0; Manatches 0; Indels 0

US-09-382-685A-2

LRNGTH: 21

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Match 100.0%; Score 108; DB 17; Length 21; Local Similarity 100.0%; Pred. No. 7.3e-10. Tidels 0 less 20; Conservative 0; Mismatches 0; Indels 0

1 VPOSIDWRDYGAVTSVKNON 20

1997-02-25 MBER: GB 9703827.7

1997-02-25

SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 3 PRIOR FILING DATE: 1997-0: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1997-0: NUMBER OF SEQ 1D NOS: 5

. Ananas comosus

US-09-382-688-3

Query Match Best Local S ORGANISM Matches

PRIOR APPLICATION NUMBER: GB 9"
PRIOR FILING DATE: 1997-02-28
PRIOR APPLICATION NUMBER: GB 9" APPLICATION NUMBER: CH FILING PATE

GB 9706119.6 1997-03-25 MBER: GB 9704252.7 GB 9703850.9

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; MOLECULE TYPE: protein
US-08-244-796-15
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US-09-791-537-84882
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CURRENT APPLICATION NUMBER: US/09/791,537
CURRENT FILLEG DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 13955
200FWARE: Patentin vetsion 3.0
SEQ ID NO 8899
LENGTH. 3667
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 15:
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TELECOMMUNICATION INFORMATION
TELEPHONE: (1516) 742 4343
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TYPE: amino acid
STRANDEDNESS: single
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       100.0%;
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       Score 108; DB 21; Length 356;
Pred. No. 1.6e-08;
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       US-09-791-537-69324; Sequence 69324, Application US/09791537; GENERAL INFORMATION:
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GENERAL INFORMATION:
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SEQ ID NO 20
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Best Local Similarity 90.0%; Pred. No. 4.1e-07;
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CURRENT FILING DATE: 2010-09-01
PRIOR APPLICATION NUMBER: PCT/GB99/02784
PRIOR FILING DATE: 1999-08-24
PRIOR FILING DATE: 1999-08-24
PRIOR APPLICATION NUMBER: GB 9819138
PRIOR APPLICATION NUMBER: GB 9819138
PRIOR FILING DATE: 1998-09-02
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APPLICANT: MYMOSE, Tracey LEHAMME
APPLICANT: GOSGESET, BEN
TITUE OF INCENTION: NUCLEIC ACIDS AND PROTEINS FROM DINEAPPLE STEM
FILE REFERENCE, 0623,090000
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ORGANISM: Ananas comosus
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; TYPE: PRT
; ORGANISM: Ananas comosus
US-09-743-987-16
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GENERAL INFORMATION:
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SEQ ID NO 16
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APPLICANT, Crossett. BEN
TITE OF INVENTION NUCLEE (ALTON AND PROTEINS FROM EINEAPEL STEW
CREENST PARLICATION NUMBER: US/00/741/987
CURRENT FILLION DES. 2001.0-01.7
BELOR APPLICATION NUMBER: DET/0889/02784
BELOR APPLICATION NUMBER: DET/0889/02784
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BELOR MERICATION NUMBER: DET/0889/02784
BELOR MERICATION NUMBER: DET/0889/02784
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APPLICANT: Crossett, Ben
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                                                        Local Similarity
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Pred. No. 4e-07;
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90.0%; Pred. No. 4.5e-07;
tive 0; Mismatches 2; Indels
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HALOR PELLON NUMBER: PCT/GESS/00550
PROOF PELLON DATE: 1299-70-23 9796119.6
PROOF PELLON NUMBER: GE 979422.7
PROOF PELLON NUMBER: 1099-102-38 979422.7
PROOF PELLON NUMBER: 1099-102-38 9791360.0
PROOF PELLON DATE: 1399-1-72-38 9791360.0
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FILE REFERENCE, 0623 079001
CHEMINET PROLICE NUMBER: US 09/202, 685A
PRICES PROLICETOR NUMBER: US 09/202, 685A
PRICES PROLICETOR NUMBER: US 09/202, 685A
PRICES PROLICETOR NUMBER: 08 070213.6
PRICES PROLICETOR NUMBER: 08 070213.6
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1997-02-25
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GENERAL INFORMATION:
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APPLICANT: Engwerda, Christian
APPLICANT: Peek, Keith
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Best Local Similarity 90.08
Matches 18, Conservative
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SEQ ID NO 4
         ; ORGANISM: Ananas comosus
US-09-791-537-84889
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PRIOR FILING DATE: 1997-0
NUMBER OF SEQ ID NOS: 5
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APPLICANT: Debe, DEPURING TO THE STRUCTURES OF PROTEIN PARLIESS AND FAMILY NEBBE TITLE OF INVESTIGATION THREE DIMENSIONAL STRUCTURES OF PROTEIN PARLIESS AND FAMILY NEBBE TITLE OF INVESTIGATION REPORTS OF THE STRUCTURES AND FAMILY NEBBE TITLE OF THE STRUCTURES OF THE STRUCTURES AND FAMILY OF THE STRUCTURES OF THE STRUCTURES OF THE STRUCTURE STRUCTURES OF THE STRUCTURE STRUCTURES OF THE STRUCTURE STRUCTURES OF THE STRUCTURES
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PITTLE OF INVENTION: THREED DIMENSIONAL STRUCTURES OF PROTEIN FAMILIES AND FAMILY MEMGE
TITTLE OF INVENTION: METHODS OF USE THEREOF
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90.0%; Pred. No. 4.1e-07;
Live 0; Mismatches 2; Indels
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CHERENT FILMS DATE: 2011.08-01
CHERENT FILMS DATE: 3011.08-01
PRIOR FILMS DATE: 1999-08-24
PRIOR FILMS DATE: 1999-08-24
PRIOR FILMS DATE: 1999-08-24
STARRER OF BOX DATE: 1998-09-02
SOFTHARE: PRIORITY NAMES OF SECULOR STARRER OF SECULOR NOS: 7.0
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CURRENT FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 153055
SOFTWARE: Faternin version 3.0
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J Sequence 84889, Application US/09791537

, GENERAL INFORMATION:
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GENERAL INFORMATION:
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SEQ ID NO 69324
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Sequence 5. Napitastico Us/995010A.
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US-09-750-210A-5
Search completed: November 21, 2002, 16:15:42 
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Best Local Similarity 94.7%; Pred. No. 4.2e-08;
Natches 18; Conservative 1; Mismatches 0; Indels
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LENGTH: 20
TYPE: PRT
ORGANISM: Ananas comosus
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Best Local Similarity 94.7%; Pred. No. 4.2e.08;
Matches 18; Conservative 1; Mismatches 0; Indels
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ALIGNMENTS	990-004A-1	-09-327-373-165	-09-791-537-13	-10-219-220-16	-09-990-004A-1	-09-327-373-16	-10-219-220	-09-990-004A-3	-09-327-373-17	-60-360-039-642	-09-791-537-3491	-09-791-537-682	-09-791-537-5675	-09-515-038-6	-60-324-109-2919	-10-219-999-6086	-09-791-537-9	-09-791-537-8488	-10-143-650-	-09-791-537-8	-09-519-283-4	-10-219-999-	-09-791-537-8488	9-743-987-	-09-791-537-6	3-5	-09-743-987-16	-09-791-537-8471	-09-791-537-3225	9-791-537-8	S-09-791-537-8487	08-244-796-1	-09-750-210	-09-382-688-	ò	-09-674-738	-09-750-210A	9-382-688-5	-09-382-6857	
	e 165	equence 165, Ap	equence 13319	equence 167,	nce 167,	e 167,	1Ce 176	e 176,	e 176,	e 6416,	e 34918,	equence 68257,	e 56752,	equence 69, App	e 29193,	e 60863,	e 91358,	e 84884,	e 19, App	e 82405,	4, Appl	e 45992,	nce 84889,	nce 14, App	nce 69324,	Sequence 20, Appl	nce 16, Apr	nce 84711,	32256	equence 84882,	ce 84879,	ce 15,	e 3, Appl	e 3, Appl	equence 2, Appl	equence 1, Appl	equence 5, Appl	equence 5,	equence 4, Appl	

US-09-382-688-4

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100.04; Score 105; DB 21; Length 216;
Bet Local 25:Milarity 100.04; Pred. No. 4.9e-08
Matches 20; Conservative 0; Mismatches 0; Indels 0;
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100.0%; Score 105; DB 21; Length 20;
Best Local Similarity 100.0%; Pred, No. 3.7e-09.
Makethes 20; Conservative 0; Mismatches 0; Indels of
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APPLICANT: Engwerder, Christian
APPLICANT: Peek, Keith
THILE OF INVENTION: Component of Bromelain
FILE REPRENCE: 0623,0740001
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CURRENT FILING DATE: 2000-12-29
PRIOR PILING DATE: US 09/382,688
PRIOR FILING DATE: 1999-08-25
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PRIOR FLING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: GB 9706119.6
PRIOR FLING DATE: 1997-03-25
PRIOR APPLICATION NUMBER: GB 97052.7
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PRIOR FILING DATE: 1997-02-25
NUMBER OF SEQ ID NOS: 5
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GENERAL INFORMATION:
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, ORGANISM: ORGANISM: Ananas comosus
US-09-750-210A-4
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GENERAL INFORMATION:
          1 VPOSIDWRDSGAVTSVKNOG 20
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Sequence 2. Application up/056/138

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1. Similarity 100.0%; Pred. Mo. 3.7e.09;
20; Conservative 0; Mismatches n: Trahir n
Query Match 100.0%; Score 105; DB 17; Length 20; DB est Local Similarity 100.0%; Pred. no. 3.7e-03. Matches 20; Conservative 0; Mismatches 0; Indels (
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Deet Local Similarity 10.04; Pred, No. 3.-6-01; 
Matches 20; Conservative 0; Mismatches 0; Indels 0
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FILE REPRENCE, 1673,704001.

CHERRY PREVIOUS NUMBER, 1059,9142,684A

CHERRY FILING DATE, 1359,01-55,504,01659

FROM FILING DATE, 1359,01-55,704,01659

FROM FILING NUMBER, 03 704119,6

FROM FILING PARTE, 1397,01-57,014,0166

FROM FILING NUMBER, 03 704122,7

FROM FILING DATE, 1397,02-25

FROM FILING DATE, 1397,02-25

FROM FILING DATE, 1397,02-25

FROM FILING DATE, 1397,02-25

FROM FILING DATE, 1397,02-25
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GENERAL INFORMATION:
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US-09-674-738-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mynott, Tracy Lehanne
APPLICANT: Engwerda, Christian
APPLICANT: Engwerda, Christian
APPLICANT: Peek, Keith
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NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                  1 VPQSIDWRDSGAVTSVKNQG 20
                                                                                                                                                                                                                                                            1 VPOSIDWRDSGAVTSVKNQG 20
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RESULT 3 US-09-674-738-2

US-09-382-688-4 LENGTH: 20 SEQ ID NO 4 TYPE: PRT Query Match

Marches

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Gaps

RESULT 6

1 VPQSIDWRDSGAVTSVKOVQC 20

TYPE: PRT LENGTH:

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CORRENT FILMS DATE: 1999-06-25
PRICE APPLICATION NUMBER: PC/1889/0093
PRICE PLILMS DATE: 1998-02-25
PRICE PLILMS DATE: 1998-02-25
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PRICE PLILMS DATE: 1998-02-25
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                                                                                                                                                                                                                                                                                                       В
                                                                                             Sequence 5, Application US/0938268BA, GENERAL INFORMATION:
                                                                                                                                                                US-09-382-688-5
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GENERAL INFORMATION:
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LENGTH: 345
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APPLICANT:
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APPLICANT, BADOMAK, Inc.
APPLICANT, BADOMAK, INc.
APPLICANT, BADOMAK, BADOMAK, STRUCTURES OF PROTEIN FAMILES AND FAMILY MEMBER
TITLE OF INVENTIOR, METHODO OF USE THEREOF
PILE REPERBOCE, FALL/JOHN, METHODO OF USE THEREOF
CILE REPERBOCE, FALL/JOHN, METHOD OF USE
CHEMBER FILMS INC.
CHEMBER FILMS INC.
METHOD FILMS
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CURRENT FILING DATE: 1999-08-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Component of Bromelain
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Mynott, Tracy Lehanne
Engwerda, Christian
Peek, Keith
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100.0%; Pred. No. 8.2e-08;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90.0%; Pred. No. 2.1e-07; Length 20;
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; ORGANISM: Ananas comosus
US-09-382-688-5
      US-09-674-738-1
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Best Local Similarity
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PRIOR FILING DATE: 1997-02-25
PRIOR APPLICATION NUMBER: GB
PRIOR FILING DATE: 1997-02-25
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN Ver: 2.0
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PRIOR APPLICATION NUMBER: GB 9
PRIOR FILING DATE: 1997-02-28
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PRIOR FILING DATE: 1998-0
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1997-02-25
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90.0%; Pred. No. 2.1e-07;
tive 1; Mismatches 1; Indels
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Gaps

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Sequence 5, Application US/09750210A
GENERAL INFORMATION:
APPLICANT: Mymott, Tracy Lehanne
APPLICANT: Engwerda, Christian
APPLICANT: Eeek, Keith
                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/750,210A
CURRENT FILING DATE: 200012-29
PRIOR APPLICATION NUMBER: US 09/382,688
PRIOR FILING DATE: 1939-08-25
PRIOR FILING DATE: 1939-08-25
PRIOR FILING DATE: 1939-08-25
PRIOR APPLICATION NUMBER: PCT/GB98/00590
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FILE REFERENCE: 0623.0740001
CURRENT APPLICATION NUMBER: US/09/382,688A
CURRENT FILING DATE: 1999-08-25
PRIOR FILING DATE: 1997-02-28
PRIOR APPLICATION NUMBER: GB 9703850.9
PRIOR FILING DATE: 1997-02-25
PRIOR APPLICATION NUMBER: GB 9703827.7
                                                                                                                                 PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: GB 9706119.6
PRIOR FILING DATE: 1997-03-25
                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Component of Bromelain
FILE REFERENCE: 0623.0740001
                                                                                                      PRIOR APPLICATION NUMBER: GB 9704252.7
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10MBER: GB_9706119.6
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                                              CORREST FILMS DATE: 1999-10-28

CORREST FILMS DATE: 1999-10-28

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FILMS AFFLORM NUMBER: 02 37-018-6, 10-28

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US-08-244-796-15
; Sequence 15, Application US/08244796
                       FILE REFERENCE: 0623.0740001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VPQSIDWRDSGAVTSVKNQ 19
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TYPE: PRT
OKGANISM: Ananas comosus
US-09-382-688-3
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US-09-750-210A-3
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Sequence 1, Application US/95/4738
ORNEAL, INFOWNEROM A.TORNIMITED GRAPHAM A.TORNIMITED GRAPHAM T.TORNIMITED OF A.TORNIMITED GRAPHAM T.TORNIMITED OF A.TORNIMITED GRAPHAM T.TORNIMITED GRAPHAM T.T. OF THORNIMITED OF A.TORNIMITE OF THORNION TO CARGINATION TITLE OF THORNION TO CARGINATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.6%; Score 93; DB 17; Length 21; 94.7%; Pred. No. 3.2e-07; tive 0; Mismatches 1; Indels
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PRIOR FILMO MRYER: 1997-02-25
PRIOR APPLICATION NUMBER: GB 9704252.7
PRIOR FILING DATE: 1997-02-28
PRIOR FILING DATE: 1997-02-25
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PRIOR FILING DATE: 1997-02-25
NUMBER OF SEQ ID NOS: 4
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GENERAL INFORMATION:
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US-09-382-685A-2
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US-09-791-537-84879
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US-08-244-796-15
                                                                                                                                                                                                                                                                                                                               APRICANT: Cabe, Derek
APRICANT: Cabe, Derek
APRICANT: Cabe, Carek
APRICANT: Cabe, Carek
TILL OF LAWRENCH ARE ILLEGATE OF PROTEIN FAMILIES AND FAMILY APRICA
TILL OF LAWRENCH ARE CABE, CABE, CABE, THERROP
FILE REFERENCE: SLIVIO 050 188 THERROP
FILE REFER
            Matches 18; Conserv
                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 84879, Application US/09791537
GENERAL INFORMATION:
APPLICANT: Bionomix, Inc.
APPLICANT: Debe, Derek
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Best Local Similarity
Matches 18; Conserv
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PILING DATE: 15-MG-1994
PAPOLICATION NAMBER: US PAPOLICATION NAMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (1516) 742 4343
TELEPHA: (1516) 742 4366
INFORMATION FOR SEQ ID NO: 15:
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ZIP: 11530-0299
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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ADDRESSEE: SCULLY SCOTT MURPHY & PRESSER
STREET: 400 GARDEN CITY PLAZA, GARDEN CITY
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LENGTH: 212 amino acids
TYPE: amino acid
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STRANDEDNESS:
TOPOLOGY: lin
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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REFERENCE/DOCKET NUMBER: 9316
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COUNTRY: UNITED STATES OF AMERICA
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ch 88.6%; Score 93; DB 21; Length 356;
11 Similarity 94.7%; pred, No. 7e-06;
18; Conservative 0; Mismatches 1; Indels
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ilarity 94.7%; Pred. No. 4e-06;
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Search completed: November 21, 2002, 16:15:43
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Sequence 80521, A
Sequence 80531, A
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THE REPRESENTE: 128 14.4 Compage
THE REPRESENTE: 128 14.4 Compage
COMMENT FILMO LTD: 2000-1328/724.676
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UNBERT FILMO LTD: 2000-1328/724.876
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SOFTWARE: Patentin version 1.2
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GENERAL INFORMATION:
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GENERAL INFORMATION:
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63.2%; Pred. No. 0.00045;
                                                                                                       64.8%; Score 70; DB 5; Length 230
63.2%; Pred. No. 0.00045;
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Sequence 80621, Application US/09724676 GENERAL INFORMATION:

US-09-724-676-80621

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Pred. No. 0.00045,
4, Mismatches 3; Indels
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Ouery Match

64.8%; Score 70; DB 5; Length 230;

Ber Local Similarity 63.2%; Pred. No. 0.00045;

Matches 12; Conservative 4; Mismatches 3; Indels
63.2%; Pred. No. 0.00045;
tive 4: Mismatches 3: Indele
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Sequence vieto, Application UN/09724676
APPLICATION COMPANION.
THE REPRESENCE 1.22814 Companion Companion Companion Application NAMES: 180.724,676
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SEQUENCE PROPERTY NAME
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NUMBER TERM DATE 200: 11.28
NUMBER SERVICE AND AUTHORIZE PROBLEM AND SERVICE AND SER
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; Sequence 80627, Application US/09724676
; GENERAL INFORMATION:
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GENERAL INFORMATION:
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Best Local Similarity 63.2
Matches 12, Conservative
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US-09-724-676-80625
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US-09-724-676-80626
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Best Local Similarity
Matches 12, Conservi
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US-09-724-676-80625
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63.2%; Pred. No. 0.00045;
Live 4; Mismatches 3; Indels
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63.2%, Pred. No. 0.00045;
tive 4; Mismatches 3, Indels
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APPRICARY, Compagn LT, Compagn

US-09-724-676-80622 ; Sequence 80622, Application US/09724676 ; GENERAL INFORMATION:

RESULT 4 ò

115 LPRSVDWREKGYVTPVKNQ 133

1 VPOSIDWRDYGAVTSVKNO 19

12; Conservative

Query Match Best Local Similarity Matches 12; Conserv

TYPE: PRT ORGANISM: Homo sapiens US-09-724-676-80621

LENGTH

Sequence 8064, Application US/09714676

APPLOAD TO ADMINISTRATION:
THE MORE MANAGEMENT AND THE THE SET OF THE

, ORGANISM: Homo sapiens US-09-724-676-80624

LENGTH: 230

TYPE: PRT Ouery Match

115 LPRSVDWREKGYVIPVKNO 133

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RESULT 5 US-09-724-676-80624

1 VPOSIDWRDYGAVTSVKNQ 19

63.2%;

Best Local Similarity 63.2 Matches 12; Conservative

Query Match

TYPE: PRT CRGANISM: Homo sapiens US-09-724-676-80622

SEQ ID NO 80622

LENGTH .

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US-09-724-676-80628
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US-09-724-676-80629
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GRHERAL MEROMATICAN:
TITLE 0F INVESTION: Variants of alternative splicing
TITLE 0F INVESTION NAMESE: 10/9/174.676
CURRENT PLILAG DATE: 2000-11-28
NAMESES OF SEQ 110 NAMES: 9722.
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CURRENT APPLICATION NAMBERS: 189/99/724.676
CURRENT FILLING DATE: 2000-11-28
NUMBER OF SEQ 10 DOS: 97222
SOPTWARE: Patentin Version 3.2
SEQ ID NO 80629
LENGTH: 230
TYPE: PRT
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GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative aplicing
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                                                                                                                                        Sequence 80630, Application US/09724676
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Best Local Similarity
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APPLICANT: Compusen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129:141.4 Compusen
CURRENT APPLICATION NUMBER: US/09/724,676
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Pred. No. 0.00045;
4; Mismatches 3
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Pred. No. 0.00045;
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Pred. No. 0.0045;
4; Mismatches 3; Indels
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US-09-724-676-80630
                           US-09-724-676-80588
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FILE REFERENCE: 12314.4 Company

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CRESH'S PERIODE 
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CURBENT APPLICATION INMERS: U8/09/724.676
CURBENT FILING DATE: 2000-11-28
NUMBER 0P 580 ID NOS: 97222
SOPTWARE: PACENTIN VETSION 3.2
SQ ID NO 86631
LENGTH: 230
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GENERAL INFORMATION:
APPLICANT: Compugen LTD
TITLE OF INVENTION: Variance of alternative splicing
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NUMBER OF SEQ ID NOS: 97222
SOFTWARE: Patentin version 3.2
SEQ ID NO 80630
LENGTH: 230
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     sequence 80588, Application US/09724676
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TYPE: PRT
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                                                                                                                                              115 LPRSYDWREKGYVTPVKNQ 133
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63.2%; Pred. No. 0.00045
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Pred. No. 0.00045;
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Query Match 64.8%; Score 70; DB 5; Length 250; Best Local Similarity 63.2%; Fred. No. 0.00018. Mexiches 12; Conservative 4; Mismatches 3; Indels
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ORGANISM: Homo sapiens

US-09-724-676-80588
SENERAL INFORMATION:
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Db 115 LPRSVDWREKGYVTPVKNQ 133

1 VPQSIDWRDYGAVTSVKNQ 19

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RESULT 15 SEGURE OF THE SEGURE

TYPE: PRT ORGANISM: Homo sapiens US-09-724-676-80589

Query Match 64.8%; Score 70; DB 5; Length 250; Beer Local Similarity 63.2%; Pred. No. 0.0000, Marches 12; Conservative 4; Mismatches 3; Indels Marches 12; Conservative 4; Mismatches 3; Indels 1 VPQSIDWRDYGAVTSVKNQ 19 :|:|:|||| | || |||| 115 LPRSVDWREKGYVTPVKNQ 133 ò

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Search completed: November 21, 2002, 16:16:04 Job time : 8 secs

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2002 Compugen Ltd
9 US-09-950-064-2
10 US-09-956-8
11 US-10-114-64-8
12 US-10-114-64-8
13 US-10-114-64-7
12 US-10-114-64-7
10 US-09-93-956-2
12 US-10-114-64-2
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10 US-09-935-300-1693
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Sequence 3, Appli
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, SEO ID NO 3
; LENGTH: 21
; TYPS: PRT
; ORGANISM: Ananas comosus
US-09-750-210A-3
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APPLICANT: Engwerds, Cl
APPLICANT: Peek, Keith
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38	3.8	38	3.8	3.8	39	39.5	40	40	40	40	40	40	40	40	40	40.5	40.5	40.5	41	41	4.	42	42	49.5	49.5
35.2	5	35.2	ů,	35.2	36.1	36.6	.7	37.0		37.0	7	37.0	.7	7	37.0	37.5	37.5	7	38.0	8	8	œ	38.9	5	45.8
518	457	420	0	307	394	165	1247	1247	1172	1172	1168	637	637	571	68	837	906	806	565	224	185	425	26	364	335
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e 12		11260,		Sequence 20, Appl	Sequence 12007, A		Sequence 13841, A	Sequence 10145, A	Sequence 2, Appli		Sequence 2, Appli	Sequence 2, Appli	.55	Sequence 1031, Ap		e 454, Ap	Sequence 1118, Ap	•	e 13	e 16	e 78	Sequence 14, Appl	*	Sequence 478, App	Sequence 10, Appl

ALIGNMENTS

PRIOR PAPICATION MANAGE: US 09/124.688
PRODR FILMO DATE: 1399 052
PRODR PRODRE MANAGE: 059 052.7
PRODR FILMO DATE: 1399 052.8
PRODR FILMO DATE: 1399 052.8 Seguence 3, Application US/09750210A Patent No. US20020102253A1 TITUE OF INVENTION: Component of Bromelain FILE REFERENCE: 0623.074001 CURRENT APPLICATION UNMBER: US/09/750,210A CURRENT FILING DATE: 2000-12-29 Tracy Lehanne da, Christian

Local Similarity 1 VPQSIDWRDYGAVTSVXXQN 20 100.0%; Score 108; DB 10; iilarity 100.0%; Pred. No. 6.9e-12; Conservative 0; Mismatches 0; Length 21; Indels

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RESULT

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18-109-78-7104-5
Segment So. 18970-18010A
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ORGANISM: ORGANISM: Ananas comosus

US-09-750-210A-5 LENGTH: 20 TYPE: PRT

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Query Match Best Local Similarity Matches 18; Conserva

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Pred, No. 1.2e.08;
0; Mismatches 2; Indels
Best Local Similarity 94.7%; Pred. No. 2e-09;
Matches 18; Conservative 0; Mismatches 1; Indels
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APPLICANT: Beggerd, Christian
APPLICANT: Beek, Meith
TITES OF INNENTONIC COmponent of Bromelain
FILE REFERENCE: 0623,074001
CURRENT PALLOKAN WINDER: 180/99/780,210A
CURRENT FILING DATE: 2000-12-29
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Bepticant: Engwerde, Christian
APPLICANT: Peek, Kath
TILL OF INDSTIGN: Coeponent of Bromelain
FILE REPRENCE: 0623.0740001
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CURRENT FILING DATE: 2000-12-29
PRIOR APPLICATION NUMBER: US 09/382,688
PRIOR FILING DATE: 1999-08-25
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PRIOR FILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: PCT/GB98/00590
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PRIOR FILING DATE: 1998-02-25
PRIOR APPLICATION NUMBER: GB 9706119.6
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PRIOR APPLICATION NUMBER: GB 9704252.7
PRIOR FILING DATE: 1997-02-28
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PRIOR FILING DATE: 1997-02-25
PRIOR APPLICATION NUMBER: GB 9703827.7
PRIOR FILING DATE: 1997-02-25
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ORGANISM: ORGANISM: Ananas comosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09750210A
Patent No. US20020102253A1
GENERAL INFORMATION:
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) Sequence 1, Application US/09750210A

; Patent No. US20020102253A1
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Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                           1 VPQSIDWRDYGAVTSVXNQ 19
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US-09-750-210A-2
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1 Similarity 94.7%; Pred, No. 4.4e-10;
118; Conservative 1; Mismacches 0; Indels
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DB 10; Length 20;

86.1%; Score 93;

ORGANISM: ORGANISM: Ananas comosus

US-09-750-210A-4

ID NO 4 TYPE: PRT Ouery Match

PROON PROTICATION NAMES OF 55 0.52 (see PROON PLANCE) AND PROON PROTICATION NAMES PERFORMANCES OF PROON PLANCE NAMES OF 1994-0.25 (see PROON PLANCE) AND PROON PLANCE NAMES OF 1994-0.25 (see PROON PLANCE) AND PROON PLANCE NAMES OF 1994-0.25 (see PROON PLANCE NAMES NAMES OF 1994-0.25 (software NAMES NAME

APPLICANT: Mynort, Tracy Lehanne APPLICANT: Deservia, Christian APPLICANT: Deservia, Component of Bromelain FILE REPRENCE: 0423, 044001, 829, 929, 92104 CURRENT FALLICATION NUMBER 192, 929, 92104

Sequence 4, Application US/09750210A Patent No. US20020102253A1

SENERAL INFORMATION: US-09-750-210A-4

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TIE DEPENDENT TO COMMENT FOR USER FOR CAME TO COMMENT APPLICATION NUMBER 100/04/62.846 CURRENT PILLING DATE; 2000-00-13 PRIOR APPLICATION NUMBER 100-07-14 PRIOR APPLICATION NUMBER 100-07-14 PRIOR PILLING DATE; 1939-07-14 PRIOR PILLING DATE; 1939-07-15 PRIOR PILLING DATE; 1939-07
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TYPE: PRT
; ORGANIEM: Bacillus subtilis
US-09-462-846-3
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1
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Patent No. US20020072107A1
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Patent No. USZ00Z0031807A1
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                                                                                          Matches
                                                                                                                                                                     Query Match
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Best Local Similarity 63.2%; Pred. No. 0.00012;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: 09/219,441
PRIOR FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANY: Hastings, et al.
TITLE OF INVENTION: HUMAN OSTEOCLAST-DERIVED CATHEPSIN
FILE REFERENCE: PF107D2D:
CURRENT APPLICATION NUMBER: US/09/953,956
CURRENT ETIME NATE: 001.00.10
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TITLE OF INVENTION: Proteases From Gram-Positive Organisms
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TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                   Match 63.9%; Score 69; DB 10; Length 333; Local Similarity 66.7%; Pred. No. 0.00041;
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        2 PQSIDWRDYGAVTSVKNQ 19
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                                                                                  Conservative
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Pred. No. 9.3e-06;
2; Mismatches 4, Indels
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                                                                          3; Indels
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; Sequence 2, Application US/09990064
; Patent No. US20020164765A1
                                                                                  US-09-990-064-2
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US-09-529-063-1
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                                                                                                                                    RESULT 10
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SEQ ID NO 9
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Patent No. US20020102542A1
GENERAL INFORMATION:
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Best Local Similarity
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Patent No. US20020142448A1
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CURRENT PILING DATE: 2002-04-03
PRIDE APPLICATION NUMBER: 08/553,125
PRIDE RILING DATE: 1995-11-07
PRIDE APPLICATION NUMBER: 08/208,007
PRIDE APPLICATION NUMBER: 08/208,007
PRIDE APPLICATION NUMBER: 08/208,007
PRIDE PILING DATE: 1994-03-08
NUMBER: 0F SEG LID NOS: 13-08
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APPLICANT: SHIBAYAMA, SHIRO
APPLICANT: TADA, HIDBAKI
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                                                                                                                                                                                                                                                       114 LPKSVDWRKKGYVTPVKNQ 132
                                                                                                                                                                                                                                                                                                                                  1 VPQSIDWRDYGAVTSVKNQ 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                            62.0%; Score 67; DB 10; Length 334;
63.2%; Pred. No. 0.00088;
stive 3; Mismatches 4; Indels
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                                                                                                                  59.3%; Score 64; DB 12; Length 331;
57.9%; Pred, No. 0.0027;
tive 3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           52.8%; Score 57; DB 10; Length 329; 66.7%; Pred. No. 0.039;
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66.7%; Pred. No. 0.039;
cive 1; Mismatches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/951,956
CURRENT TELING DATE: 2001-00-18
PRIOR APPLICATION NUMBER: 09/219,441
PRIOR FARM DATE: 1398-12-23
NUMBER OF SEQ ID NOS: 14
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Patent No. US20020142448Al
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               115 LPDSVDWREKGCVTEVKYQ 133
                                                                                                                                                                                                                                                                                      1 VPOSIDWRDYGAVTSVKNO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 POSIDWRDYGAVTSVKNG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 POSIDWRDYGAVTSVKNQ 19
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                                                                                                                          Query Match 59.3%
Best Local Similarity 57.9%
Matches 11, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 66.77
Matches 12, Conservative
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Best Local Similarity 66.74
Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.1
, ORGANISM: Homo sapiens
US-10-114-464-8
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JS-09-953-956-7
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JS-10-114-464-7
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TYPE: PRT
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US-10-114-464-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59.3%; Score 64, DB 9, Length 331, 57.9%; Pred. No. 0.0027; tive 3; Mismatches 5; Indels
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PROFER NO. USSODOIL4446A1
REPURENT INFORMATION:
APPLICATION OF THE SET OF 
                                PARTICANT FAVALURES TO PROTECT OR THE STREET OF THE STREET
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CURRENT APPLICATION NUMBER: US/10/114,464
CURRENT FILING DATE: 2002-04-03
PRIGR FULLIGATION WORBER: 08/533,125
PRIGR FILING DATE: 1995-11-07
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Matches 11; Conservative
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CRGANISM: Homo sapiens
US-09-953-956-8
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        GENERAL INFORMATION
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US-10-114-464-8
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LENGTH: 331
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score greater than or equal to the score of the result being printed, 
and is derived by analysis of the total score distribution.
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1 / (902) Z / (robota // 2)/ (robota // 2)

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US-09-750-210A-5
US-09-750-210A-2
US-09-750-210A-2
US-09-750-210A-1
US-09-750-210A-1
US-09-953-956-9
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US-09-953-956-9
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       US-10-114-464-2
US-09-925-300-1693
US-09-953-956-10
US-10-114-464-10
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                                                                                                                                                                                                                                                                                                                                                                               US-09-953-956-8
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Sequence 4, Appli
Sequence 3, Appli
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Sequence 7, Appli
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ALIGNMENTS	-09-862-007-	US-09-815-242-13686	-729-674	-09-922-217-1	US-09-922-217-1117	-729-9	-09-800-729-	-09-	US-09-986-676A-2	US-09-815-242-10370	US-09-811-284-227	-09-738-973-1	-09-	US-10-016-768-11	-10-016-	-09-909	-09-782-980	1-669-	US-09-893-737-78	242-1384	-09-815-242-	-820-893	301	US-09-864-761-42873	100	US-09-925-302-478
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	2, Appli	13686, A	160, App	1118, Ap	1117, Ap	95, Appl	130, App	<ol><li>Appli</li></ol>			227, App	101, App	100, App	11, Appl	10, Appl	<ol><li>Appli</li></ol>	15, Appl	412, App	78, Appl	13841, A		50, Appl	1302, Ap	2	1021, Ap	478, App

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SEQ ID NO 4

LENGTH: 20

TYPE: PRT

ORGANISM: ORGANISM:
US-09-750-210A-4
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US-09-750-210A-4
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RECOR APPLICATION NOWERS US 90/18-588
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Patent No. US20020102253A1
GENERAL INFORMATION:
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Engwerda, Ch
APPLICANT: Peek, Keith
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SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: GB 9703827.7
PRIOR FILING DATE: 1997-02-25
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                                                                                  Length 20;
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Matches 1 VPQSIDWRDSGAVTSVKNQG 20 VPQSIDWRDSGAVTSVKNQG 20; 100.0%; Score 105; DB 10; tlarity 100.0%; Pred. No. 1.8e-11; Conservative 0; Mismatches 0; Indels

> 0; Gaps

> > 0

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RESULT

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APPLICATION TO THE SEPTION THE

Sequence 5, Application US/09750210A Patent No. US20020102253A1

US-09-750-210A-5

GENERAL INFORMATION:

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Pred. No. 3.3e-07;
0; Mismatches 3; Indels
Best Local Similarity 94.7%; Pred. No. 1.9e-09;
Matches 18; Conservative 0; Mismatches 1; Indels
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PRIOR PATRON DAYS (1959)
PRIOR PATRON
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Parblicant: Engwerde Christian
APPLICANT: Peek, Keith
TITLE OF INDENTION: Component of Bromelain
FILE REFRENCE: 0623.0740001
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CURRENT FILING DATE: 2000-12-29
PRIOR APPLICATION NUMBER: US 09/382,688
PRIOR FILING DATE: 1999-08-25
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Beplicant: Engwards, Christian
APPLICANT: Peek, Keith
THIE OF INDENTION: Component of Bromelain
FILE REFRENCE: 0623.0740001
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RAIDE APPLICATION NUMBER: US 09/382,688 RAIDE FILING DATE: 099-08-28 RAIDE FILING PRICE FILING NUMBER: PCT/6898/00590
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PRIOR APPLICATION NUMBER: GB 9706119.6
PRIOR FILING DATE: 1997-03-25
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PRIOR APPLICATION NUMBER: GB 9703850.9
PRIOR FILING DATE: 1997-02-25
PRIOR PELICATION NUMBER: GB 9703827.7
PRIOR FILING DATE: 1997-02-25
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ORGANISM: ORGANISM: Ananas comosus
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Patent No. US20020102253A1
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Patent No. US20020102253A1
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Best Local Similarity 83.3%;
Matches 15; Conservative
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Query March 89:5%; Score 94; DB 10; Length 20; Best Local Similarity 90:0%; Pred. No. 1.2e-09; Marches 18; Conservative 1; Mismatches 1; Indels Marches 18; Conservative 1; Mismatches 18; Indels

US-09-750-210A-3 ; Sequence 3, Application US/09750210A ; Patent No. US20020102253A1

SENERAL INFORMATION:

TYPE: PRT ORGANISM: ORGANISM: Ananas comosus

US-09-750-210A-5

O ID NO 5 LENGTH: 20

88.6%; Score 93; DB 10; Length 21;

TYPE: PRT ORGANISM: ORGANISM: Ananas comosus

US-09-750-210A-3

Query Match LENGTH

SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 3 NUMBER OF SEQ ID NOS: 5

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                                                                                                                  ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-953-956-9
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Best Local Similarity 70.0
Best Local Similarity 70.0
                                                                                                                                                                                             SEQ ID NO 9
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Best Local Similarity 65.0%; Pred. No. 5.5e-06;
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CURRENT PRILICATION INDEERS US/03/462, 846

CURRENT FLILED ONE: 2000-07-123

RIGH RYLLOWING MUSES US-7-125

RIGH RYLLOW INDEES US-7-15

RIGH RYLLOW INDEES US-7-15

RIGH RYLLOW INDEES US-7-15
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Patent No. US20020031807A1
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                                       Matches
                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9, Application US/09953956
Patent No. US20020072107A1
                                                                                                                                                                                                                                     FILE REFERENCE: PF107D2D1
CURRENT APELICATION NUMBER: US/09/953,956
CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: 09/219,441
PRIOR FILING DATE: 1998-12-23
NUMBER: OF SED ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Hastings, et al.
TITLE OF INVENTION: HUMAN OSTEOCLAST-DERIVED CATHEPSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                     SOFTWARE: Patentin Ver. 2.1
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ORGANISM: Bacillus subtilis
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POSIDWRDSGAVTSVKNQG 20
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                                       13; Conservative
                                                       h 73.3%; Score 77; DB 10; Length 333; Similarity 68.4%; Pred. No. 1.9e-05;
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Pred. No. 3.9e-07;
2; Mismatches 4
                                       Mismatches
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                                                                                                                                                          RESULT 10
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                   GENERAL IMPORMATION:
APPLICANT: Hastings, et al.
TITLE OF INVENTION: HUMAN OSTEOCHAST-DERIVED CATHERSIN
FILE REFERENCE: PF107D5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8, Application US/09953956
Patent No. US20020072107A1
GENERAL INFORMATION:
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Patent No. US20020142448A1
                                                                                              Sequence 8, Application US/10114464
Patent No. US20020142448A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 8
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                                                                                                                                                                                                                                                                                                    Matches
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CUURRENT FILING DATE: 2007-09-18
PRIOR APPLICATION NUMBER: 09/219,441
PRIOR FILING DATE: 1998-12-23
MUMBER 09 850 ID NOS: 14
SECTIVARS: PATENTIN VOT. 2.1
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PRIOR PILING DATE: 1995-11-07
PRIOR APPLICATION NUMBER: 08/208,007
PRIOR FILING DATE: 1994-03-08
NUMBER OF SEQ ID NOS: 14
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CURRENT FILING DATE: 2002-04-03
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   CURRENT APPLICATION NUMBER: US/10/114,464
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3; Mismatches 5
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Query Match

G6.7%; Score 69; DB 10; Length 334;
Best Local Similarity 63.4%; Pred. No. 0.0004;
Best Excellant 12; Conservative 3; Wismatches 4; Indels
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Best Local Similarity 68.4%; Pred. No. 0.0019. 
Matches 13; Conservative 1; Mismanches 5; Indels
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CURRENT APPLICATION NUMBER: 09/29.96
PRIOR APPLICATION NUMBER: 09/29,441
PRIOR APPLICATION NUMBER: 09/29,441
NUMBER OF SEQ ID NOS: 44
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; Sequence 7, Application US/09953956
; Patent No. US20020072107A1
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, Sequence 7, Application US/10114464
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ORGANISM: Homo sapiens
            ; ORGANISM: Homo sapiens
US-09-529-063-1
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TYPE: PRT
ORGANISM: Homo sapiens
US-09-953-956-7
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DB 95 Local Similarity 60.0%; Pred. no. 0.00019. 
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CURRENT FILING DATE: 2002-04-03
PRICA APPLICATION NOMBER: 04/53,125
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Patent No. US20020164765Al
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, ORGANISM: Homo sapiens
US-10-114-464-8
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US-09-990-064-2
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